## SEQUENCE LISTING



240

<110> Chappell, Joseph Noel, Joseph P. Starks, Courtney M. Manna, Kathleen R.

## <120> SYNTHASES

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			20 aat					25		_			30			1.	44
Ser	Ile	Asp	Asn	Gln	Val	Ala	Glu	Lys	Tyr	Ala	Gln	Glu	Ile	Glu	Ala	1.4	**

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gcc gat aca ttg aat ttg att gac att att gaa cgc ctt ggt ata tcc

40

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caa Gln	aac Asn	tca Ser	aac Asn 100	cya Cya	aat Asn	gat Asp	ttg Leu	tgc Cys 105	acc Thr	tct Ser	gca Ala	ctt Leu	caa Gln 110	ttt Phe	cga Arg		336
ttg Leu	ctc Leu	agg Arg 115	caa Gln	cac His	ggt Gly	ttc Phe	aac Asn 120	atc Ile	tct Ser	cct Pro	gaa Glu	att Ile 125	ttc Phe	agc Ser	aaa Lys		384
ttc Phe	caa Gln 130	gat Asp	gaa Glu	aat Asn	ggc Gly	aaa Lys 135	ttc Phe	aag Lys	gag Glu	tct Ser	ctt Leu 140	gct Ala	agt Ser	gat Asp	gtc Val		432
tta Leu 145	gga Gly	tta Leu	tta Leu	aac Asn	ttg Leu 150	tat Tyr	gaa Glu	gct Ala	tca Ser	cat His 155	gta Val	agg Arg	act Thr	cat His	gct Ala 160		480
gac Asp	gat Asp	atc Ile	tta Leu	gaa Glu 165	gac Asp	gca Ala	ctt Leu	gct Ala	ttc Phe 170	tcc Ser	act Thr	atc Ile	cat His	ctt Leu 175	gaa Glu		528
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tct Ser	gca Ala	gct Ala	cca Pro 180	cat His	ttg Leu	aaa Lys	tct Ser	cca Pro 185	ctt Leu	agg Arg	gag Glu	caa Gln	gtg Val 190	aca Thr	cat His	,	576
Ser	gca Ala ctt Leu	Ala	Pro 180 caa	His	Leu	Lys	Ser	Pro 185 ggt	Leu	Arg	Glu	Gln	Val 190 gag	Thr	His	,	576 624
gcc Ala	Ala	gag Glu 195	Pro 180 caa Gln tca	tgt Cys tca	ttg Leu atc	Lys cac His	aag Lys 200	Pro 185 ggt Gly	gtt Val	cct Pro	Glu aga Arg	gtc Val 205	Val 190 gag Glu aat	Thr acc Thr	His cga Arg	•	
gcc Ala	Ala ctt Leu ttc Phe	gag Glu 195 atc Ile	Pro 180 caa Gln tca Ser	tgt Cys tca Ser	ttg Leu atc Ile	cac His tat Tyr 215	aag Lys 200 gac Asp	Pro 185 ggt Gly aag Lys	gtt Val gaa Glu	Arg cct Pro caa Gln	Glu aga Arg tcg ser 220	Gln gtc Val 205 aag Lys	Val 190 gag Glu aat Asn	Thr acc Thr aat Asn	His cga Arg gtg Val	•	624
gcc Ala ttc Phe tta Leu 225	Ala ctt Leu ttc Phe 210 ctt	gag Glu 195 atc Ile cga Arg	Pro 180 caa Gln tca Ser ttt Phe	tgt Cys tca Ser gcc Ala	ttg Leu atc Ile aaa Lys 230 caa	cac His tat Tyr 215 ttg Leu	aag Lys 200 gac Asp gat Asp	Pro 185 ggt Gly aag Lys ttc Phe	gtt Val gaa Glu aac Asn	cct Pro caa Gln ttg Leu 235	aga Arg tcg Ser 220 ctc Leu	gtc Val 205 aag Lys cag Gln	Val 190 gag Glu aat Asn atg Met	Thr acc Thr aat Asn ttg Leu	His  Cga Arg  gtg Val  cac His 240		624
gcc Ala ttc Phe tta Leu 225 aaa Lys gta	Ala ctt Leu ttc Phe 210 ctt Leu caa	gag Glu 195 atc Ile cga Arg gaa Glu aca	Pro 180 caa Gln tca Ser ttt Phe ctt Leu ctt	tgt Cys tca Ser gcc Ala gct Ala 245	ttg Leu atc Ile aaa Lys 230 caa Gln	cac His tat Tyr 215 ttg Leu gta Val	aag Lys 200 gac Asp gat Asp tca Ser	Pro 185 ggt Gly aag Lys ttc Phe agg Arg	gtt Val gaa Glu aac Asn tgg Trp 250	cct Pro caa Gln ttg Leu 235 tgg Trp	aga Arg tcg Ser 220 ctc Leu aaa Lys	gtc Val 205 aag Lys cag Gln gat Asp	Val 190 gag Glu aat Asn atg Met ttg Leu	Thr acc Thr aat Asn ttg Leu gat Asp 255 tac	His  Cga Arg  gtg Val  cac His 240  ttt Phe		624 672 720

Met Leu Val Lys Thr Ile Ser Met Ile Ser Ile Val Asp Asp Thr Phe 290 300 300 300 300 300 300 300 300 300 3				-	-													
Asp Ala Tyr Gly Thr Val Lys Glu Leu Glu Ala Tyr Thr Asp Ala Ile 310  caa aga tgg gat atc aac gaa att gat cgg ctt cct gat tac atg aaa 10 Gln Arg Trp Asp Ile Asn Glu Ile Asp Arg Leu Pro Asp Tyr Met Lys 325  atc agt tat aaa gct att cta gat ctc tac aag gat tat gaa aag gaa Ile Ser Tyr Lys Ala Ile Leu Asp Leu Tyr Lys Asp Tyr Glu Lys Glu Arg 340  ttg tct agt gcc gga aga tct cat att gtc tgc cat gca ata gaa aga Ileu Ser Ala Gly Arg Ser His Ile Val Cys His Ala Ile Glu Arg 360  atg aaa gaa gaa gta gta aga aat tat aat gtc gag tca aca tgg ttt att Met Lys Glu Arg Asp Tyr Asn Val Glu Ser Thr Trp Phe Ile 370  gaa gga tat atg cca cct gtt tct gaa tac cta agc aat gca cta gca Glu Gly Tyr Met Pro Pro Val Ser Glu Tyr Leu Ser Asn Ala Leu Ala 390  act acc aca tat tac tac ctc gcg aca aca tcg tat ttg ggc atg aag Thr Thr Thr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys 410  tct gaa gct acg gag caa gat ttt gag tgg ttg ta tat gas acg act gca aca aca tcg Lys Ile 420  ctt gaa gct agt gta att att att att att att gcc gag tt att gag ac aca aca tcg Lys Ile 420  ctt gaa gct agt gta att att att gt cga gtt att gag ac aca aca tcg Lys Ile 420  ctt gaa gct agt gta att att att att att act gc gag tt att gag ac aca gcc acg Lys Ile Glu Ala Ser Val Ile Asp Asp Thr Ala Thr Ala Thr Glu Glu Lys Ser Afg Gly Gln Ile Asp Asp Thr Ala Thr Glu Val Glu Lys Ser Afg Gly Gln Ile Ala Thr Gly Ile Glu Cys Afg Wet Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465		Leu					Ser					Val						912
Arg Trp Asp Ile Asn Glu Ile Asp Arg Leu Pro Asp Tyr Met Lys 335  atc agt tat aaa gct att cta gat ctc tac aag gat tat gaa aag gaa lle Ser Tyr Lys Ala Ile Leu Asp Leu Tyr Lys Asp Tyr Glu Lys Glu 340  ttg tct agt gcc gga aga tct cat att gtc tgc cat gca ata gaa aga lle Ser Ser Ala Gly Arg Ser His Ile Val Cys His Ala Ile Glu Arg 355  atg aaa gaa gta gta aga aat tat aat gtc gag tca aca tgg ttt att lys Glu Gly Tyr Met Pro Pro Val Ser Glu Tyr Leu Ser Asn Ala Leu Ala 390  gaa gga tat atg cca cct gtt tct gaa tac cta agc aat gca cta gca Glu Gly Tyr Met Pro Pro Val Ser Glu Tyr Leu Ser Asn Ala Leu Ala 390  act acc aca tat tac tac ctc gcg aca aca tcg tat ttg ggc atg aag and 12 Thr Thr Thr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys 405  tct gcc acg gag caa gat ttt gag tgg ttg tca aag aat cca aaa att Ser Ala Thr Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile 420  ctt gaa gct agt gta att ata tgt cga gtt atc gat gac aca aca tcg tat leu Gly Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr Ala Thr Ala Thr Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr Ala Thr Ala Thr Glu Glu Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr Ala Thr Glu Val Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys Arg Cyal Ala Met Ala Lys Phe 450  tgc atg aga gat tat ggt ata tca aca aca aga gcc atg gca atg gca atg gca atg gca atg aga gca atg gca at	Asp	Ala '				Val					Āla					Ile		960
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Leu Ser Ser Ala Gly Arg Ser His Ile Val Cys His Ala Ile Glu Arg 355 Ala Gly Arg Ser His Ile Val Cys His Ala Ile Glu Arg 365 Ala Gly Arg 360 Ala Glu Ser Thr Trp Phe Ile 370 Ala Arg Asn Tyr Asn Val Glu Ser Thr Trp Phe Ile 370 Ala Arg Asn Tyr Asn Val Glu Ser Thr Trp Phe Ile 370 Ala Arg Asn Tyr Asn Val Glu Ser Thr Trp Phe Ile 370 Ala Arg Asn Tyr Asn Val Glu Ser Thr Trp Phe Ile 370 Ala 385 Ala Leu Ala 390 Ala Arg Asn Ala Leu Ala 395 Ala Leu Ala 395 Ala Leu Ala Thr Thr Tyr Tyr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys 410 Ala Thr Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile 420 Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr Ala Thr Ala Ser Val Ile Ile Cys Arg Val Ile Ala Thr Gly Ile Glu Cys 450 Ala Thr Gly Ile Glu Cys Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys 450 Ala Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465 Ala Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465 Ala Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465 Ala Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465 Ala Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465 Ala Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465 Ala Met Ala Lys Phe 470 Ala Ala Met Ala Lys Phe 480 Ala Met Ala Ala Met Ala Lys Phe 480 Ala Met Ala Ala Met Ala Lys Phe 480 Ala Met Ala Ala Me				Lys					Leu					Glu			. :	1056
Met Lys Glu Val Val Arg Asn Tyr Asn Val Glu Ser Thr Trp Phe Ile 370  gaa gga tat atg cca cct gtt tct gaa tac cta agc aat gca cta gca Glu Gly Tyr Met Pro Pro Val Ser Glu Tyr Leu Ser Asn Ala Leu Ala 385  act acc aca tat tac tac ctc gcg aca aca tcg tat ttg ggc atg aag Thr Thr Thr Tyr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys 410  tct gcc acg gag caa gat ttt gag tgg ttg tca aag aat cca aca act ly End Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile 425  ctt gaa gct agt gta att ata tgt cga gtt atc gat gac aca aca gc acg Leu Glu Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr 435  tac gag gtt gag aaa agc agg gga caa att gca act gga att gag tgc Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys 455  tgc atg aga gat tat ggt ata tca aca aaa gag gca atg gct aaa ttt Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 480		Ser	Ser					His					Āla				:	1104
Glu Gly Tyr Met Pro Pro Val Ser Glu Tyr Leu Ser Asn Ala Leu Ala 395  act acc aca tat tac tac ctc gcg aca aca tcg tat ttg ggc atg aag Thr Thr Thr Tyr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys 415  tct gcc acg gag caa gat ttt gag tgg ttg tca aag aat cca aaa att Ser Ala Thr Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile 420  ctt gaa gct agt gta att ata tgt cga gtt atc gat gac aca gcc acg Leu Glu Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr 435  tac gag gtt gag aaa agc agg gga caa att gca act gga att gag tgc Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys 450  tgc atg aga gat tat ggt ata tca aca aca gag gca atg gct aaa ttt Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465		Lys					Asn					Ser						1152
Thr Thr Thr Tyr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys 415  tct gcc acg gag caa gat ttt gag tgg ttg tca aag aat cca aaa att Ser Ala Thr Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile 420  ctt gaa gct agt gta att ata tgt cga gtt atc gat gac aca gcc acg Leu Glu Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr 435  tac gag gtt gag aaa agc agg gga caa att gca act gga att gag tgc Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys 455  tgc atg aga gat tat ggt ata tca aca aaa gag gca atg gct aaa ttt Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 470  12  13  14  15  16  17  18  19  19  10  10  11  12  13  14  15  16  17  18  19  19  10  10  10  11  11  12  13  14  15  16  17  18  19  19  10  10  10  11  11  12  13  14  15  16  17  18  19  19  10  10  10  10  10  10  10  10	Glu	ı Ğİy '				Pro					Leu					Ala	•	1200
Ser Ala Thr Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile 420  ctt gaa gct agt gta att ata tgt cga gtt atc gat gac aca gcc acg Leu Glu Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr 435  tac gag gtt gag aaa agc agg gga caa att gca act gga att gag tgc Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys 450  tgc atg aga gat tat ggt ata tca aca aaa gag gca atg gct aaa ttt Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465  470  425  Leu Ser Lys Asn Pro Lys Ile 430  13  14  15  16  17  18  18  19  19  10  10  11  12  13  14  15  16  17  18  19  19  10  10  10  11  11  12  13  14  15  16  17  18  18  19  19  10  10  10  10  10  10  10  10					Tyr					Thr					Met		:	1248
Leu Glu Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr 435  tac gag gtt gag aaa agc agg gga caa att gca act gga att gag tgc Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys 450  tgc atg aga gat tat ggt ata tca aca aaa gag gca atg gct aaa ttt Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465  470  480				Glu					Trp					Pro			:	1296
Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys 450 455 460  tgc atg aga gat tat ggt ata tca aca aaa gag gca atg gct aaa ttt Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465 470 475 480		ı Glu .	Āla					Cys					Asp				•	1344
Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe 465 470 475 480	tac Tyr	r Glu	gtt Val	gag Glu	aaa Lys	agc Ser	Arg	gga Gly	caa Gln	att Ile	gca Ala	Thr	gga Gly	att Ile	g <b>ag</b> Glu	tgc Cys	:	1392
cas ast ato out one are one one too ass out att ast one out off 14	Cys	s Met				Gly					Glu					Phe		1440
caa aat atg gct gag aca gca tgg aaa gat att aat gaa gga ctt ctt 14 Gln Asn Met Ala Glu Thr Ala Trp Lys Asp Ile Asn Glu Gly Leu Leu 485 490 495					Glu					Asp					Leu		;	1488

agg Arg	Pro	act Thr	Pro 500	gtc Val	tct Ser	aca Thr	gaa Glu	ttt Phe 505	tta Leu	act Thr	cct Pro	att Ile	ctc Leu 510	aat Asn	ctt Leu	<b>1</b> 536
get Ala	cgt	att Ile 515	gtt Val	gag Glu	gtt Val	aca Thr	tat Tyr 520	ata Ile	cac His	aat Asn	cta Leu	gat Asp 525	gga Gly	tac Tyr	act Thr	1584
cat His	ccq Pro 530	gag Glu	aaa Lys	gtc Val	tta Leu	aaa Lys 535	cct Pro	cac His	att Ile	act Ile	aac Asn 540	cta Leu	ctt Leu	gtg Val	gac Asp	1632
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Trp Ala Leu Gly Val Tyr Phe Glu Pro Gln Tyr Ser Gln Ala Arg Val
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Met Leu Val Lys Thr Ile Ser Met Ile Ser Ile Val Asp Asp Thr Phe
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Asp Ala Tyr Gly Thr Val Lys Glu Leu Glu Ala Tyr Thr Asp Ala Ile
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                                         315
Gln Arg Trp Asp Ile Asn Glu Ile Asp Arg Leu Pro Asp Tyr Met Lys
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Ile Ser Tyr Lys Ala Ile Leu Asp Leu Tyr Lys Asp Tyr Glu Lys Glu
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                                 345
Leu Ser Ser Ala Gly Arg Ser His Ile Val Cys His Ala Ile Glu Arg
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Met Lys Glu Val Val Arg Asn Tyr Asn Val Glu Ser Thr Trp Phe Ile
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Glu Gly Tyr Met Pro Pro Val Ser Glu Tyr Leu Ser Asn Ala Leu Ala
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Thr Thr Thr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys
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Ser Ala Thr Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile
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Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys
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Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe
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Gln Asn Met Ala Glu Thr Ala Trp Lys Asp Ile Asn Glu Gly Leu Leu
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Arg Pro Thr Pro Val Ser Thr Glu Phe Leu Thr Pro Ile Leu Asn Leu
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tc Se	c att r Ile	gat Asp 35	aat Asn	cag Gln	gtt Val	gcg Ala	gaa Glu 40	aag Lys	tat Tyr	gct Ala	caa Gln	gag Glu 45	att Ile	gaa Glu	gca Ala	144
tt Le	g aag u Lys 50	gaa Glu	caa Gln	acg Thr	agg Arg	agt Ser 55	atg Met	ctg Leu	tta Leu	gca Ala	acc Thr 60	gga Gly	agg Arg	aaa Lys	ttg Leu	192
gc Al 6	c gat a Asp 5	aca Thr	ttg Leu	aat Asn	ttg Leu 70	att Ile	gac Asp	att Ile	att Ile	gaa Glu 75	cgc Arg	ctt Leu	ggt Gly	ata Ile	tcc Ser 80	240
ta Ty	c cac r His	ttt Phe	gag Glu	aaa Lys 85	gaa Glu	att Ile	gat Asp	gag Glu	att Ile 90	ttg Leu	gat Asp	cag Gln	att Ile	tac Tyr 95	aac Asn	288
ca Gl:	a aac n Asn	tca Ser	aac Asn 100	tgc Cys	aat Asn	gat Asp	ttg Leu	tgc Cys 105	acc Thr	tct Ser	gca Ala	ctt Leu	caa Gln 110	ttt Phe	cga Arg	336
t te	g ctc ı Leu	agg Arg 115	caa Gln	cac His	ggt Gly	ttc Phe	aac Asn 120	atc Ile	tct Ser	cct Pro	g <b>aa</b> Glu	att Ile 125	ttc Phe	agc Ser	aaa Lys	384
Pho	c caa e Gln 130	gat Asp	g <b>aa</b> Glu	aat Asn	ggc Gly	aaa Lys 135	ttc Phe	aag Lys	gag Glu	tct Ser	ctt Leu 140	gct Ala	agt Ser	gat Asp	gtc Val	432
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	ttc Phe															672

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aaa Lys	caa Gln	g <b>aa</b> Glu	ctt Leu	gct Ala 245	caa Gln	gta Val	tca Ser	agg Arg	tgg Trp 250	tgg Trp	aaa Lys	gat Asp	ttg Leu	gat Asp 255	ttt Phe	768
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gag Glu	gca Ala	tta Leu 275	gga Gly	gtt Val	tat Tyr	ttt Phe	gag Glu 280	cct Pro	caa Gln	tac Tyr	tct Ser	caa Gln 285	gct Ala	cgc Arg	gtc Val	864
atg Met	ctc Leu 290	gtt Val	aag Lys	acc Thr	ata Ile	tca Ser 295	atg Met	att Ile	tcg Ser	att Ile	gtc Val 300	gat Asp	gac Asp	acc Thr	ttt Phe	912
gat Asp 305	gct Ala	tac Tyr	ggt Gly	aca Thr	gtt Val 310	aaa Lys	gaa Glu	ctt Leu	gag Glu	gca Ala 315	tac Tyr	aca Thr	gat Asp	gcc Ala	ata Ile 320	960
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cat His	ccg Pro 530	gag Glu	aaa Lys	gtc Val	tta Leu	aaa Lys 535	cct Pro	cac His	att Ile	att Ile	aac Asn 540	cta Leu	ctt Leu	gtg Val	gac Asp	• 1632	2
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 Ser

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 Asp
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Leu Leu Arg Gln His Gly Phe Asn Ile Ser Pro Glu Ile Phe Ser Lys Phe Gln Asp Glu Asn Gly Lys Phe Lys Glu Ser Leu Ala Ser Asp Val Leu Gly Leu Leu Asn Leu Tyr Glu Ala Ser His Val Arg Thr His Ala 1.55 Asp Asp Ile Leu Glu Asp Ala Leu Ala Phe Ser Thr Ile His Leu Glu Ser Ala Ala Pro His Leu Lys Ser Pro Leu Arg Glu Gln Val Thr His Ala Leu Glu Gln Cys Leu His Lys Gly Val Pro Arg Val Glu Thr Arg Fhe Phe Ile Ser Ser Ile Tyr Asp Lys Glu Gln Ser Lys Asn Asn Val Lau Leu Arg Phe Ala Lys Leu Asp Phe Asn Leu Leu Gln Met Leu His Lys Gln Glu Leu Ala Gln Val Ser Arg Trp Trp Lys Asp Leu Asp Phe Val Thr Thr Leu Pro Tyr Ala Arg Asp Arg Val Val Glu Cys Tyr Phe Glu Ala Leu Gly Val Tyr Phe Glu Pro Gln Tyr Ser Gln Ala Arg Val Met Leu Val Lys Thr Ile Ser Met Ile Ser Ile Val Asp Asp Thr Phe Asp Ala Tyr Gly Thr Val Lys Glu Leu Glu Ala Tyr Thr Asp Ala Ile Gln Arg Trp Asp Ile Asn Glu Ile Asp Arg Leu Pro Asp Tyr Met Lys Ile Ser Tyr Lys Ala Ile Leu Asp Leu Tyr Lys Asp Tyr Glu Lys Glu Leu Ser Ser Ala Gly Arg Ser His Ile Val Cys His Ala Ile Glu Arg Met Lys Glu Val Val Arg Asn Tyr Asn Val Glu Ser Thr Trp Phe Ile Glu Gly Tyr Met Pro Pro Val Ser Glu Tyr Leu Ser Asn Ala Leu Ala Thr Thr Thr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys Ser Ala Thr Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile Leu Glu Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe ~ 470 Gln Asn Met Ala Glu Thr Ala Trp Lys Asp Ile Asn Glu Gly Leu Leu Arg Pro Thr Pro Val Ser Thr Glu Phe Leu Thr Pro Ile Leu Asn Leu Ala Arg Ile Val Glu Val Thr Tyr Ile His Asn Leu Asp Gly Tyr Thr 

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                                                                       96
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                                                                      144
Ser Ile Asp Asn Gln Val Ala Glu Lys Tyr Ala Gln Glu Ile Glu Ala
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ttg aag gaa caa acg agg agt atg ctg tta gca acc gga agg aaa ttg
                                                                      192
Leu Lys Glu Gln Thr Arg Ser Met Leu Leu Ala Thr Gly Arg Lys Leu
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gcc gat aca ttg aat ttg att gac att att gaa cgc ctt ggt ata tcc
                                                                      240
Ala Asp Thr Leu Asn Leu Ile Asp Ile Ile Glu Arg Leu Gly Ile Ser
tac cac ttt gag aaa gaa att gat gag att ttg gat cag att tac aac
                                                                      288
Tyr His Phe Glu Lys Glu Ile Asp Glu Ile Leu Asp Gln Ile Tyr Asn
caa aac tca aac tgc aat gat ttg tgc acc tct gca ctt caa ttt cqa
                                                                      336
Gln Asn Ser Asn Cys Asn Asp Leu Cys Thr Ser Ala Leu Gln Phe Arq
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                                                                      384
Leu Leu Arg Gln His Gly Phe Asn Ile Ser Pro Glu Ile Phe Ser Lys
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tto caa gat gaa aat ggc aaa tto aag gag tot ott got agt gat gto
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Phe Gln Asp Glu Asn Gly Lys Phe Lys Glu Ser Leu Ala Ser Asp Val
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tta gga tta tta aac ttg tat gaa gct tca cat gta agg act cat gct
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Leu Gly Leu Leu Asn Leu Tyr Glu Ala Ser His Val Arg Thr His Ala
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gcc Ala	ctt Leu	gag Glu 195	caa Gln	C <b>ys</b>	ttg Leu	cac His	aag Lys 200	ggt Gly	gtt Val	cct Pro	aga Arg	gtc Val 205	gag Glu	acc Thr	cga Arg	624
ttc Phe	ttc Phe 210	atc Ile	tca Ser	tca Ser	atc Ile	tat Tyr 215	gac Asp	aag Lys	gaa Glu	caa Gln	tcg Ser 220	aag Lys	aat Asn	aat Asn	gtg Val	672
tta Leu 225	ctt Leu	cga Arg	ttt Phe	gcc Ala	aaa Lys 230	ttg Leu	gat Asp	ttc Phe	aac Asn	ttg Leu 235	ctc Leu	cag Gln	atg Met	ttg Leu	cac His 240	720
aaa Lys	caa Gln	gaa Glu	ctt Leu	gct Ala 245	caa Gln	gta Val	tca Ser	agg Arg	tgg Trp 250	tgg Trp	aaa Lys	gat Asp	ttg Leu	gat Asp 255	ttt Phe	768
gta Val	aca Thr	aca Thr	ctt Leu 260	cca Pro	tat Tyr	gct Ala	aga Arg	gat Asp 265	cga Arg	gta Val	gtt Val	gaa Glu	tgc Cys 270	tac Tyr	ttt Phe	816
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caa Gln	aga Arg	tg <b>g</b> Trp	gat Asp	atc Ile 325	aac Asn	gaa Glu	att Ile	gat Asp	cgg Arg 330	ctt Leu	cct Pro	gat Asp	tac Tyr	atg Met 335	aaa Lys	1008
atc Ile	agt Ser	tat Tyr	aaa Lys 340	get Ala	att Ile	cta Leu	gat Asp	ctc Leu 345	tac Tyr	aag Lys	gat Asp	tat Tyr	gaa Glu 350	aag Lys	gaa Glu	1056
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gaa Glu 385	gga Gly	tat Tyr	atg Met	cca Pro	cct Pro 390	gtt Val	tct Ser	gaa Glu	tac Tyr	cta Leu 395	agc Ser	aat Asn	gca Ala	cta Leu	gca Ala 400		1200
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tct Ser	gcc Ala	acg Thr	gag Glu 420	caa Gln	gat Asp	ttt Phe	gag Glu	tgg Trp 425	ttg Leu	tca Ser	aag Lys	aat Asn	cca Pro 430	aaa Lys	att Ile		1296
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caa Gln	aat Asn	atg Met	gct Ala	gag Glu 485	aca Thr	gca Ala	tgg Trp	aaa Lys	gat Asp 490	att Ile	aat Asn	gaa Glu	g <b>ga</b> Gly	ctt Leu 495	ctt Leu		1488
agg Arg	ccc Pro	act Thr	ccc Pro 500	gtc Val	tct Ser	aca Thr	gaa Glu	ttt Phe 505	tta Leu	act Thr	cct Pro	att Ile	ctc Leu 510	aat Asn	ctt Leu		1536
gct Ala	cgt Arg	att Ile 515	gtt Val	gag Glu	gtt Val	aca Thr	ttc Phe 520	ata Ile	cac His	aat Asn	cta Leu	gat Asp 525	gga Gly	tac Tyr	act Thr		1584
cat His	ccg Pro 530	Glu	aaa Lys	gtc Val	tta Leu	aaa Lys 535	cct Pro	cac His	att Ile	att Ile	aac Asn 540	cta Leu	ctt Leu	gtg Val	gac Asp		1632
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                                     410
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Leu Glu Ala Ser Val Ile Ile Cys Arg Val Ile Asp Asp Thr Ala Thr
Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys
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Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe
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Gln Asn Met Ala Glu Thr Ala Trp Lys Asp Ile Asn Glu Gly Leu Leu
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Arg Pro Thr Pro Val Ser Thr Glu Phe Leu Thr Pro Ile Leu Asn Leu
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Ala Arg Ile Val Glu Val Thr Phe Ile His Asn Leu Asp Gly Tyr Thr
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Val Ala Asp Phe Ser Pro Ser Leu Trp Gly Asp Gln Phe Leu Ser Phe
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tcc att gat aat cag gtt gcg gaa aag tat gct caa gag att gaa gca
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Ser Ile Asp Asn Gln Val Ala Glu Lys Tyr Ala Gln Glu Ile Glu Ala
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Leu Lys Glu Gln Thr Arg Ser Met Leu Leu Ala Thr Gly Arg Lys Leu
gcc gat aca ttg aat ttg att gac att att gaa cgc ctt ggt ata tcc
                                                                      240
Ala Asp Thr Leu Asn Leu Ile Asp Ile Ile Glu Arg Leu Gly Ile Ser
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Tyr His Phe Glu Lys Glu Ile Asp Glu Ile Leu Asp Gln Ile Tyr Asn
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t to Let	g ctc 1 Leu	agg Arg 115	caa Gln	cac His	ggt Gly	ttc Phe	aac Asn 120	atc Ile	tct Ser	cct Pro	gaa Glu	att Ile 125	ttc Phe	agc Ser	aaa Lys		394
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Gln Arg Trp Asp Ile Asn Glu Ile Asp Arg Leu Pro Asp Tyr Met Lys
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Ile Ser Tyr Lys Ala Ile Leu Asp Leu Tyr Lys Asp Tyr Glu Lys Glu
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Leu Ser Ser Ala Gly Arg Ser His Ile Val Cys His Ala Ile Glu Arg
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                                                 525
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Ser Ile Lys Ile
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                                                                       48
Met Ala Ser Ala Ala Val Ala Asn Tyr Glu Glu Glu Ile Val Arg Pro
gte gee gae tte tee eet agt ete tgg ggt gat eag tte ett tea tte
                                                                       96
Val Ala Asp Phe Ser Pro Ser Leu Trp Gly Asp Gln Phe Leu Ser Phe
```

			-	•													
tcc Ser	att Ile	gat Asp 35	aat Asn	cag Gln	gtt Val	gcg Ala	gaa Glu 40	aag Lys	tat Tyr	gct Ala	caa Gln	gag Glu 45	att Ile	gaa Glu	gca Ala		144
	aag Lys 50																192
	gat Asp																240
tac Tyr	cac His	ttt Phe	gag Glu	aaa Lys 85	gaa Glu	att Ile	gat Asp	gag Glu	att Ile 90	ttg Leu	gat Asp	cag Gln	att Ile	tac Tyr 95	aac Asn		288
caa Gln	aac Asn	tca Ser	aac Asn 100	tgc Cys	aat Asn	gat Asp	ttg Leu	tgc Cys 105	acc Thr	tct Ser	gca Ala	ctt Leu	caa Gln 110	ttt Phe	cga Arg		336
	ctc Leu															٠	384
ttc Phe	caa Gln 130	gat Asp	gaa Glu	aat Asn	ggc Gly	aaa Lys 135	ttc Phe	aag L <b>ys</b>	gag Glu	tct Ser	ctt Leu 140	gct Ala	agt Ser	gat Asp	gtc Val	•	432
	gga Gly																480
	gat Asp																528
tct Ser	gca Ala	gct Ala	cca Pro 180	cat His	ttg Leu	aaa Lys	tct Ser	cca Pro 185	ctt Leu	agg Arg	gag Glu	caa Gln	gtg Val 190	aca Thr	cat His		576
	ctt Leu															-	624
	ttc Phe 210																672
tta Leu 225	ctt Leu	cga Arg	ttt Phe	gcc Ala	aaa Lys 230	ttg Leu	gat Asp	ttc Phe	aac Asn	ttg Leu 235	ctc Leu	cag Gln	atg Met	t t g Leu	cac His 240		720

aaa Lys	caa Gln	gaa Glu	ctt. Leu	gct Ala 245	caa Gln	gta Val	tca Ser	agg Arg	tgg Trp 250	tgg Trp	aaa Lys	gat Asp	ttg Leu	gat Asp 255	ttt Phe	768
gta Val	aca Thr	aca Thr	ctt Leu 260	cca Pro	tat Tyr	gct Ala	aga Arg	gat Asp 265	cga Arg	gta Val	gtt Val	gaa Glu	tgc Cys 270	tac Tyr	ttt Phe	816
tcg Ser	gca Ala	tta Leu 275	gga Gly	gtt Val	tat Tyr	ttt Phe	gag Glu 280	cct Pro	caa Gln	tac Tyr	tct Ser	caa Gln 285	gct Ala	cgc Arg	gtc Val	864
atg Met	ctc Leu 290	gtt Val	aag Lys	acc Thr	ata Ile	tca Ser 295	atg Met	att Ile	tcg Ser	att Ile	gt.c Val 300	gat Asp	gac Asp	acc Thr	ttt Phe	912
gat Asp 305	gct Ala	tac Tyr	ggt Gly	aca Thr	gtt Val 310	aaa Lys	gaa Glu	ctt Leu	gag Glu	gca Ala 315	tac Tyr	aca Thr	gat Asp	gcc Ala	ata Ile 320	960
caa Gln	aga Arg	tgg Trp	gat Asp	atc Ile 325	aac Asn	gaa Glu	att Ile	gat Asp	cgg Arg 330	ctt Leu	cct Pro	gat Asp	tac Tyr	atg Met 335	aaa Lys	1008
atc Ile	agt Ser	tat Tyr	aaa Lys 340	gct Ala	att Ile	cta Leu	gat Asp	ctc Leu 345	tac Tyr	aag Lys	gat Asp	tat Tyr	gaa Glu 350	aag Lys	gaa Glu	. 1056
ttg Leu	tct Ser	agt Ser 355	gcc Ala	gga Gly	aga Arg	tct Ser	cat His 360	att Ile	gtc Val	tgc Cys	cat His	gca Ala 365	ata Ile	gaa Glu	aga Arg	1104
atg Met	aaa Lys 370	gaa Glu	gta Val	gta Val	aga Arg	aat Asn 375	tat Tyr	aat Asn	gtc Val	gag Glu	tca Ser 380	aca Thr	tgg Trp	ttt Phe	att Ile	1152
gaa Glu 385	gga Gly	tat Tyr	atg Met	cca Pro	cct Pro 390	gtt Val	tct Ser	gaa Glu	tac Tyr	cta Leu 395	agc Ser	aat Asn	gca Ala	cta Leu	gca Ala 400	1200
act Thr	acc Thr	aca Thr	tat Tyr	tac Tyr 405	tac Tyr	ctc Leu	gcg Ala	aca Thr	aca Thr 410	tcg Ser	tat Tyr	ttg Leu	ggc Gly	atg Met 415	aag Lys	1248
tct Ser	gcc Ala	acg Thr	gag Glu 420	caa Gln	gat Asp	ttt Phe	gag Glu	tgg Trp 425	ttg Leu	tca Ser	aag Lys	aat Asn	cca Pro 430	aaa Lys	att Ile	1296
ctt Leu	gaa Glu	gct Ala 435	agt Ser	gta Val	att Ile	ata Ile	tgg Trp 440	cga Arg	gtt Val	atc Ile	gat Asp	gac Asp 445	aca Thr	gcc Ala	acg Thr	1344

tac Tyr	gag Glu 450	gtt Val	gag Glu-	aaa Lys	agc Ser	agg Arg 455	gga Gly	caa Gln	att Ile	gca Ala	act Thr 460	gga Gly	att Ile	gag Glu	tgc Cys	1392
tgc Cys 465	atg Met	aga Arg	gat Asp	tat Tyr	ggt Gly 470	ata Ile	tca Ser	aca Thr	aaa Lys	gag Glu 475	gca Ala	atg Met	gct Ala	aaa Lys	ttt Phe 480	1440
caa Gln	aat Asn	atg Met	gct Ala	gag Glu 485	aca Thr	gca Ala	tgg Trp	aaa Lys	gat Asp 490	att Ile	aat Asn	gaa Glu	gga Gly	ctt Leu 495	ctt Leu	1488
agg Arg	ccc Pro	act Thr	ccc Pro 500	gtc Val	tct Ser	aca Thr	gaa Glu	ttt Phe 505	tta Leu	act Thr	cct Pro	att Ile	ctc Leu 510	aat Asn	ctt Leu	1536
gct Ala	cgt Arg	att Ile 515	gtt Val	gag Glu	gtt Val	aca Thr	tat Tyr 520	ata Ile	cac His	aat Asn	cta Leu	gat Asp 525	gga Gly	tac Tyr	act Thr	1584
cat His	ccg Pro 530	gag Glu	aaa Lys	gtc Val	tta Leu	aaa Lys 535	cct Pro	cac His	att Ile	att Ile	aac Asn 540	cta Leu	ctt Leu	gtg Val	gac Asp	1632
		aaa Lys		·												1644

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<213> TEAS W273S/C440W

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Leu Gly Leu Leu Asn Leu Tyr Glu Ala Ser His Val Arg Thr His Ala
                    150
                                         155
Asp Asp Ile Leu Glu Asp Ala Leu Ala Phe Ser Thr Ile His Leu Glu
                165
                                    170
Ser Ala Ala Pro His Leu Lys Ser Pro Leu Arg Glu Gln Val Thr His
                                185
Ala Leu Glu Gln Cys Leu His Lys Gly Val Pro Arg Val Glu Thr Arg
                            200
Phe Phe Ile Ser Ser Ile Tyr Asp Lys Glu Gln Ser Lys Asn Asn Val
                        215
                                             220
Leu Leu Arg Phe Ala Lys Leu Asp Phe Asn Leu Leu Gln Met Leu His
                    230
                                        235
Lys Gln Glu Leu Ala Gln Val Ser Arg Trp Trp Lys Asp Leu Asp Phe
                                    250
Val Thr Thr Leu Pro Tyr Ala Arg Asp Arg Val Val Glu Cys Tyr Phe
            260
                                265
Ser Ala Leu Gly Val Tyr Phe Glu Pro Gln Tyr Ser Gln Ala Arg Val
                            280
                                                 285
Met Leu Val Lys Thr Ile Ser Met Ile Ser Ile Val Asp Asp Thr Phe
    290
                        295
                                            300
Asp Ala Tyr Gly Thr Val Lys Glu Leu Glu Ala Tyr Thr Asp Ala Ile
                    310
                                        315
Gln Arg Trp Asp Ile Asn Glu Ile Asp Arg Leu Pro Asp Tyr Met Lys
                325
                                    330
Ile Ser Tyr Lys Ala Ile Leu Asp Leu Tyr Lys Asp Tyr Glu Lys Glu
            340
                                345
Leu Ser Ser Ala Gly Arg Ser His Ile Val Cys His Ala Ile Glu Arg
                            360
Met Lys Glu Val Val Arg Asn Tyr Asn Val Glu Ser Thr Trp Phe Ile
                        375
Glu Gly Tyr Met Pro Pro Val Ser Glu Tyr Leu Ser Asn Ala Leu Ala
                    390
                                        395
Thr Thr Thr Tyr Tyr Leu Ala Thr Thr Ser Tyr Leu Gly Met Lys
                405
                                    410
Ser Ala Thr Glu Gln Asp Phe Glu Trp Leu Ser Lys Asn Pro Lys Ile
                                425
                                                     430
Leu Glu Ala Ser Val Ile Ile Trp Arg Val Ile Asp Asp Thr Ala Thr
                            440
        435
                                                445
Tyr Glu Val Glu Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys
                        455
Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe
                    470
                                        475
Gln Asn Met Ala Glu Thr Ala Trp Lys Asp Ile Asn Glu Gly Leu Leu
                                    490
Arg Pro Thr Pro Val Ser Thr Glu Phe Leu Thr Pro Ile Leu Asn Leu
            500
                                505
Ala Arg Ile Val Glu Val Thr Tyr Ile His Asn Leu Asp Gly Tyr Thr
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His Pro Glu Lys Val Leu Lys Pro His Ile Ile Asn Leu Leu Val Asp
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Ser Ile Lys Ile
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gtc gcc gac ttc tcc cct agt ctc tgg ggt gat cag ttc ctt tca ttc
                                                                       96
Val Ala Asp Phe Ser Pro Ser Leu Trp Gly Asp Gln Phe Leu Ser Phe
tcc att gat aat cag gtt gcg gaa aag tat gct caa gag att gaa gca
                                                                      144
Ser Ile Asp Asn Gln Val Ala Glu Lys Tyr Ala Gln Glu Ile Glu Āļa
         35
ttg aag gaa caa acg agg agt atg ctg tta gca acc gga agg aaa ttg
                                                                      192
Leu Lys Glu Gln Thr Arg Ser Met Leu Leu Ala Thr Gly Arg Lys Leu
     50
gcc gat aca ttg aat ttg att gac att att gaa cgc ctt ggt ata tcc
                                                                      240
Ala Asp Thr Leu Asn Leu Ile Asp Ile Ile Glu Arg Leu Gly Ile Ser
                     70
tac cac ttt gag aaa gaa att gat gag att ttg gat cag att tac aac
                                                                      288
Tyr His Phe Glu Lys Glu Ile Asp Glu Ile Leu Asp Gln Ile Tyr Asn
                 85
                                      90
caa aac tca aac tgc aat gat ttg tgc acc tct gca ctt caa ttt cga
                                                                      336
Gln Asn Ser Asn Cys Asn Asp Leu Cys Thr Ser Ala Leu Gln Phe Arg
                                 105
ttg ctc agg caa cac ggt ttc aac atc tct cct gaa att ttc agc aaa
                                                                      384
Leu Leu Arg Gln His Gly Phe Asn Ile Ser Pro Glu Ile Phe Ser Lys
        115
                             120
tto caa gat gaa aat ggc aaa tto aag gag tot ott got agt gat gto
                                                                       432
Phe Gln Asp Glu Asn Gly Lys Phe Lys Glu Ser Leu Ala Ser Asp Val
    130
                         135
tta qqa tta tta aac ttq tat qaa qct tca cat qta aqq act cat qct
                                                                       480
Leu Gly Leu Leu Asn Leu Tyr Glu Ala Ser His Val Arg Thr His Ala
145
                     150
                                         155
                                                             160
gac gat atc tta gaa gac gca ctt gct ttc tcc act atc cat ctt gaa
                                                                       528
Asp Asp Ile Leu Glu Asp Ala Leu Ala Phe Ser Thr Ile His Leu Glu
                165
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tct Ser	gca Ala	gct Ala	cca Pro 180	cat His	ttg Leu	aaa Lys	tct Ser	cca Pro 185	ctt Leu	agg Arg	gag Glu	caa Gln	gtg Val 190	aca Thr	cat His		576
	ctt Leu																624
	ttc Phe 210																672
	ctt Leu																720
aaa Lys	caa Gln	gaa Glu	ctt Leu	gct Ala 245	caa Gln	gta Val	tca Ser	agg Arg	tgg Trp 250	tgg Trp	aaa Lys	gat Asp	ttg Leu	gat Asp 255	ttt Phe		768
gta Val	aca Thr	aca Thr	ctt Leu 260	cca Pro	tat Tyr	gct Ala	aga Arg	gat Asp 265	cga Arg	gta Val	gtt Val	gaa Glu	tgc Cys 270	tac Tyr	ttt Phe	٠	816
	gca Ala															•	864
	ctc Leu 290																912
	gct Ala																960
	aga Arg																1008
	agt Ser																1056
	tct Ser																1104
atg Met	aaa Lys 370	gaa Glu	gta Val	gta Val	aga Arg	aat Asn 375	tat Tyr	aat Asn	gtc Val	gag Glu	tca Ser 380	aca Thr	tgg Trp	ttt Phe	att Ile		1152

gaa Glu 385	gga Gly	tat Tyr	atg Met	cca Pro	cct Pro 390	gtt Val	tct Ser	gaa Glu	tac Tyr	cta Leu 395	agc Ser	aat Asn	gca Ala	cta Leu	gca Ala 400		1200
act Thr	acc Thr	aca Thr	tat Tyr	tac Tyr 405	nns Xaa	nns Xaa	gcg Ala	aca Thr	aca Thr 410	tcg Ser	tat Tyr	ttg Leu	ggc Gly	atg Met 415	aag Lys		1248
tct Ser	gcc Ala	acg Thr	gag Glu 420	caa Gln	gat Asp	ttt Phe	gag Glu	tgg Irp 425	ttg Leu	tca Ser	aag Lys	aat Asn	cca Pro 430	aaa Lys	att Ile		1296
ctt Leu	gaa Glu	gct Ala 435	agt Ser	gta Val	att Ile	ata Ile	tgt Cys 440	cga Arg	gtt Val	atc Ile	gat Asp	gac Asp 445	aca Thr	gcc Ala	acg Thr		1344
tac Tyr	gag Glu 450	gtt Val	gag Glu	aaa Lys	agc Ser	agg Arg 455	gga Gly	caa Gln	att Ile	gca Ala	act Thr 460	gga Gly	att Ile	gag Glu	tgc Cys		1392
tgc Cys 465	atg Met	aga Arg	gat Asp	tat Tyr	ggt Gly 470	ata Ile	tca Ser	aca Thr	aaa Lys	gag Glu 475	gca Ala	atg Met	gct Ala	aaa Lys	ttt Phe 480	-	1440
			gct Ala														1488
agg Arg	ccc Pro	act Thr	ccc Pro 500	gtc Val	tct Ser	aca Thr	gaa Glu	ttt Phe 505	tta Leu	act Thr	cct Pro	att Ile	ctc Leu 510	aat Asn	ctt Leu		1536
gct Ala	cgt Arg	att Ile 515	gtt Val	gag Glu	gtt Val	aca Thr	tat Tyr 520	ata Ile	cac His	aat Asn	cta Leu	gat Asp 525	gga Gly	tac Tyr	act Thr		1584
			aaa Lys														1632
	atc																1644

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440
Tyr Glu Val Glu-Lys Ser Arg Gly Gln Ile Ala Thr Gly Ile Glu Cys
                        455
                                             460
Cys Met Arg Asp Tyr Gly Ile Ser Thr Lys Glu Ala Met Ala Lys Phe
                    470
                                         475
Gln Asn Met Ala Glu Thr Ala Trp Lys Asp Ile Asn Glu Gly Leu Leu
                485
                                    490
Arg Pro Thr Pro Val Ser Thr Glu Phe Leu Thr Pro Ile Leu Asn Leu
            500
                                 505
                                                     510
Ala Arg Ile Val Glu Val Thr Tyr Ile His Asn Leu Asp Gly Tyr Thr
                             520
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His Pro Glu Lys Val Leu Lys Pro His Ile Ile Asn Leu Leu Val Asp
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Ser Ile Lys Ile
545
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catgcccaaa tacgatgttg tcgcsnnsnn gtaatatgtg gtagttgcta gtgc
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      <220>
      <221> CDS
      <222> (6)...(1892)
      <223> pinene synthase
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cagea atg get eta-gtt tet ace gea eeg ttg get tee aaa tea tge etg
                                                                        50
      Met Ala Leu Val Ser Thr Ala Pro Leu Ala Ser Lys Ser Cys Leu
cac aaa teg ttg atc agt tet acc cat gag ett aag get ete tet aga
                                                                        98
His Lys Ser Leu Ile Ser Ser Thr His Glu Leu Lys Ala Leu Ser Arg
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aca Thr	att Ile	cca Pro	gct Ala 35	cta Leu	gga Gly	atg Met	agt Ser	agg Arg 40	cga Arg	Gly 999	aaa Lys	tct Ser	atc Ile 45	act Thr	cct Pro		146
tcc Ser	atc Ile	agc Ser 50	atg Met	agc Ser	tct Ser	acc Thr	acc Thr 55	gtt Val	gta Val	acc Thr	gat Asp	gat Asp 60	ggt Gly	gta Val	cga Arg		194
aga Arg	cgc Arg 65	atg Met	ggc	gat Asp	ttc Phe	cat His 70	tcc Ser	aac Asn	ctc Leu	tgg Trp	gac Asp 75	gat A <b>sp</b>	gat Asp	gte Val	ata [le		242
cag Gln 80	tct Ser	tta Leu	cca Pro	acg Thr	gct Ala 85	tat Tyr	gag Glu	gaa Glu	aaa Lys	tcg Ser 90	tac Tyr	ctg Leu	gag Glu	cgt Arg	gct Ala 95		290
gag Glu	aaa Lys	ctg Leu	atc Ile	999 Gly 100	gaa Glu	gta Val	aag Lys	aac Asn	atg Met 105	ttc Phe	aat Asn	tcg Ser	atg Met	tca Ser 110	tta Leu		338
gaa Glu	gat Asp	gga Gly	gag Glu 115	tta Leu	atg Met	agt Ser	ccg Pro	ctc Leu 120	aat Asn	gat Asp	ctc Leu	att Ile	caa Gln 125	cgc Arg	ctt Leu	٠	386
tgg Trp	att Ile	gtc Val 130	gac Asp	agc Ser	ctt Leu	gaa Glu	cgt Arg 135	ttg Leu	gly aaa	atc Ile	cat His	aga Arg 140	cat His	ttc Phe	aaa Lys	•	434
gat Asp	gag Glu 145	ata Ile	aaa Lys	tcg Ser	gcg Ala	ctt Leu 150	gat Asp	tat Tyr	gtt Val	tac Tyr	agt Ser 155	tat Tyr	tgg Trp	ggc Gly	gaa Glu		482
					999 Gly 165												530
					cga Arg												578
					gct Ala												626
					aca Thr												674
					att Ile												722

gct Ala 240	gaa Glu	atc Ile	ttc Phe	tct Ser	acc Thr 245	aaa Lys	tat Tyr	tta Leu	aaa Lys	gaa Glu 250	Ala	ctg Leu	caa Gln	aag Lys	att Ile 255	7	70
ccg Pro	gtc Val	tcc Ser	agt Ser	ctt Leu 260	tcg Ser	cga Arg	gag Glu	atc Ile	999 Gly 265	gac Asp	gtt Val	ttg Leu	gaa Glu	tat Tyr 270	Gly	81	18
tgg Trp	cac His	aca Thr	tat Tyr 275	ttg Leu	ccg Pro	cga Arg	ttg Leu	gaa Glu 280	gca Ala	agg Arg	aat Asn	tac Tyr	atc Ile 285	Gln	gtc Val	86	56
ttt Phe	gga Gly	cag Gln 290	gac Asp	act Thr	gag Glu	aac Asn	acg Thr 295	aag Lys	tca Ser	tat Tyr	gtg Val	aag Lys 300	agc Ser	aaa Lys	aaa Lys	91	L <b>4</b>
ctt Leu	tta Leu 305	gaa Glu	ctc Leu	gca Ala	aaa Lys	ttg Leu 310	gag Glu	ttc Phe	aac Asn	atc Ile	ttt Phe 315	caa Gln	tcc Ser	tta Leu	caa Gln	96	52
aag Lys 320	agg Arg	gag Glu	tta Leu	gaa Glu	agt Ser 325	ctg Leu	gtc Val	aga Arg	tgg Trp	tgg Trp 330	aaa Lys	gaa Glu	tcg Ser	ggt Gly	ttt Phe 335	101	. 0
cct Pro	gag Glu	atg Met	acc Thr	ttc Phe 340	tgc Cys	cga Arg	cat His	cgt Arg	cac His 345	gtg Val	gaa Glu	tac Tyr	tac Tyr	act Thr 350	ttg Leu	105	8
gct Ala	tcc Ser	tgc Cys	att Ile 355	gcg Ala	ttc Phe	gag Glu	cct Pro	caa Gln 360	cat His	tct Ser	gga Gly	ttc Phe	aga Arg 365	ctc Leu	ggc Gly	110	6
ttt Phe	gcc Ala	aag Lys 370	acg Thr	tgt Cys	cat His	ctt Leu	atc Ile 375	acg Thr	gtt Val	ctt Leu	gac Asp	gat Asp 38Q	atg Met	tac Tyr	gac Asp	115	4
acc Thr	ttc Phe 385	ggc Gly	aca Thr	gta Val	gac Asp	gag Glu 390	ctg Leu	gaa Glu	ctc Leu	ttc Phe	aca Thr 395	gcg Ala	aca Thr	atg Met	aag Lys	120	2
aga Arg 400	tgg Trp	gat Asp	ccg Pro	tcc Ser	tcg Ser 405	ata Ile	gat Asp	tgc Cys	ctt Leu	cca Pro 410	gaa Glu	tat Tyr	atg Met	aaa Lys	gga Gly 415	125	0
gtg Val	tac Tyr	ata Ile	gcg Ala	gtt Val 420	tac Tyr	gac Asp	acc Thr	gta Val	aat Asn 425	gaa Glu	atg Met	gct Ala	cga Arg	gag Glu 430	gca Ala	129	8
gag Glu	gag Glu	gct Ala	caa Gln 435	ggc Gly	cga Arg	gat Asp	acg Thr	ctc Leu 440	aca Thr	tat Tyr	gct Ala	cgg Arg	gaa Glu 445	gct Ala	tgg Trp	134	6
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Glu	Ala	Tyr 450	Ile	Asp	Ser	Tyr	Met 455	Gln	Glu	Ala	Arg	Trp 460	Ile	Ala	Thr	
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gac Asp	atc Ile	gcc Ala	aga Arg 595	gct Ala	ttc Phe	cat His	tac Tyr	ggc Gly 600	tac Tyr	aaa Lys	tac Tyr	cga Arg	gac Asp 605	ggc Gly	tac Tyr	1826
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gaa Glu	tct Ser 625	gtg Val	cct Pro	ttg Leu	tag *	caac	agct	ca a	atct	atgo	ec et	atgo	ctato	ī		1922
tege	gtta aaag	aa a ytt g	tata gtaat	tgtg	g aa aa aa	iggta iaaaa	gccg aaaa	g tto aaa	gatg laaa	jtag	agga	taag	rtt t	gtta	itaa <b>tt</b>	1982 2018

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Tyr Leu Pro Ser Phe Asp Glu Tyr Tyr Glu Asn Gly Lys Val Ser Cys
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Gly His Arg Ile Ser Ala Leu Gln Pro Ile Leu Thr Met Asp Ile Pro
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                          15
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Ser His Phe Lys Ser Ser Pro Lys Leu Leu Ser Ser Thr Asn Ser Ser
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			ggc Gly														772

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Cys Ser Ser Ser-Gln Leu Thr Thr Glu Arg Arg Ser Gly Asn Tyr Asn Pro Ser Arg Trp Asp Val Asn Phe Ile Gln Ser Leu Leu Ser Asp Tyr Lys Glu Asp Lys His Val Ile Arg Ala Ser Glu Leu Val Thr Leu Val Lys Met Glu Leu Glu Lys Glu Thr Asp Gln Ile Arg Gln Leu Glu Leu Ile Asp Asp Leu Gln Arg Met Gly Leu Ser Asp His Phe Gln Asn Glu Phe Lys Glu Ile Leu Ser Ser Ile Tyr Leu Asp His His Tyr Tyr Lys Asn Pro Phe Pro Lys Glu Glu Arg Asp Leu Tyr Ser Thr Ser Leu Ala Phe Arg Leu Leu Arg Glu His Gly Phe Gln Val Ala Gln Glu Val Phe Asp Ser Phe Lys Asn Glu Glu Gly Glu Phe Lys Glu Ser Leu Ser Asp Asp Thr Arg Gly Leu Leu Gln Leu Tyr Glu Ala Ser Phe Leu Leu Thr Glu Gly Glu Thr Thr Leu Glu Ser Ala Arg Glu Phe Ala Thr Lys Phe Leu Glu Glu Lys Val Asn Glu Gly Gly Val Asp Gly Asp Leu Leu Thr Arg Ile Ala Tyr Ser Leu Asp Ile Pro Leu His Trp Arg Ile Lys Arg Pro Asn Ala Pro Val Trp Ile Glu Trp Tyr Arg Lys Arg Pro Asp Met Asn Pro Val Val Leu Glu Leu Ala Ile Leu Asp Leu Asn Ile Val Gln Ala Gln Phe Gln Glu Glu Leu Lys Glu Ser Phe Arg Trp Trp Arg Asn Thr Gly Phe Val Glu Lys Leu Pro Phe Ala Arg Asp Arg Leu Val Glu Cys Tyr Phe Trp Asn Thr Gly Ile Ile Glu Pro Arg Gln His Ala Ser Ala Arg Ile Met Met Gly Lys Val Asn Ala Leu Ile Thr Val Ile Asp Asp Ile Tyr Asp Val Tyr Gly Thr Leu Glu Glu Leu Glu Gln Phe Thr Asp Leu Ile Arg Arg Trp Asp Ile Asn Ser Ile Asp Gln Leu Pro Asp Tyr Met Gln Leu Cys Phe Leu Ala Leu Asn Asn Phe Val Asp Asp Thr Ser Tyr Asp Val Met Lys Glu Lys Gly Val Asn Val Ile Pro Tyr Leu Arg Gln Ser Trp Val Asp Leu Ala Asp Lys Tyr Met Val Glu Ala Arg Trp Phe Tyr Gly Gly His Lys Pro Ser Leu Glu Glu Tyr Leu Glu Asn Ser Trp Gln Ser Ile Ser Gly Pro Cys Met Leu Thr His Ile Phe Phe 

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						acc Thr										147
tca Ser	cta Leu	caa Gln	atg Met	ggt Gly 50	aat Asn	gag Glu	atc Ile	caa Gln	act Thr 55	gga Gly	cga Arg	cga Arg	act Thr	gga Gly 60	ggc Gly	199
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	gag Glu																435
	gat Asp																483
	caa Gln																531
	gac Asp 175															-	579
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cta Leu 270	gaa Glu	cta Leu	gct Ala	aaa L <del>y</del> s	ttg Leu 275	gac Asp	ttc Phe	aat Asn	atg Met	gtt Val 280	caa Gln	gca Ala	caa Gln	ttt Phe	caa Gln 285		867
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					aaa Lys											1683
					ata Ile											1731
					aac Asn											.1779
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Asp Phe Ser Val Ser Gln Glu Val Phe Asp Cys Phe Lys Asn Asp Lys -165 Gly Thr Asp Phe Lys Pro Ser Leu Val Asp Asp Thr Arg Gly Leu Leu Gln Leu Tyr Glu Ala Ser Phe Leu Ser Ala Gln Gly Glu Glu Thr Leu His Leu Ala Arg Asp Phe Ala Thr Lys Phe Leu His Lys Arg Val Leu Val Asp Lys Asp Ile Asn Leu Leu Ser Ser Ile Glu Arg Ala Leu Glu Leu Pro Thr His Trp Arg Val Gln Met Pro Asn Ala Arg Ser Phe Ile Asp Ala Tyr Lys Arg Arg Pro Asp Met Asn Pro Thr Val Leu Glu Leu Ala Lys Leu Asp Phe Asn Met Val Gln Ala Gln Fhe Gln Glu Leu Lys Glu Ala Ser Arg Trp Trp Asn Ser Thr Gly Leu Val His Glu Leu Pro Phe Val Arg Asp Arg Ile Val Glu Cys Tyr Tyr Trp Thr Thr Gly Val Val Glu Arg Arg Glu His Gly Tyr Glu Arg Ile Met Leu Thr Lys Ile Asn Ala Leu Val Thr Thr Ile Asp Asp Val Phe Asp Ile Tyr Gly Thr Leu Glu Glu Leu Gln Leu Phe Thr Thr Ala Ile Gln Arg Trp Asp Ile Glu Ser Met Lys Gln Leu Pro Pro Tyr Met Gln Ile Cys Tyr Leu Ala Leu Phe Asn Phe Val Asn Glu Met Ala Tyr Asp Thr Leu Arg Asp Lys Gly Phe Asn Ser Thr Pro Tyr Leu Arg Lys Ala Trp Val Asp Leu Val Glu Ser Tyr Leu Ile Glu Ala Lys Trp Tyr Tyr Met Gly His Lys Pro Ser Leu Glu Glu Tyr Met Lys Asn Ser Trp Ile Ser Ile Gly Gly Ile Pro Ile Leu Ser His Leu Phe Phe Arg Leu Thr Asp Ser Ile Glu Glu Glu Asp Ala Glu Ser Met His Lys Tyr His Asp Ile Val Arg Ala Ser Cys Thr Ile Leu Arg Leu Ala Asp Asp Met Gly Thr Ser Leu Asp Glu Val Glu Arg Gly Asp Val Pro Lys Ser Val Gln Cys Tyr Met Asn Glu Lys Asn Ala Ser Glu Glu Glu Ala Arg Glu His Val Arg Ser Leu Ile Asp Gln Thr Trp Lys Met Met Asn Lys Glu Met Met Thr Ser Ser Phe Ser Lys Tyr Phe Val Gln Val Ser Ala Asn Leu Ala Arg Met Ala Gln Trp Ile Tyr Gln His Glu Ser Asp Gly Phe Gly Met Gln His Ser Leu Val Asn Lys Met Leu Arg Gly Leu Leu Phe Asp Arg Tyr Glu

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ctt gtc cct Leu Val Pro 30	tgc act gca ccc Cys Thr Ala Pro	c acc gct cgc ctc c Thr Ala Arg Leu 40	cgg gca tct tgc Arg Ala Ser Cys	tcc · 145 Ser 45
tca aaa cta Ser Lys Leu	caa gaa gct ca Gln Glu Ala Hi 50	t caa atc cga cga s Gln Ile Arg Arg 55	tct gga aac tac Ser Gly Asn Tyr 60	caa . 193 Gln
cct gcc ctt Pro Ala Leu	tgg gat tcc aa Trp Asp Ser Ass 65	t tac att cag tct n Tyr Ile Gln Ser 70	ctc aat act cca Leu Asn Thr Pro 75	tat 241 Tyr
acg gag gag Thr Glu Glu 80	agg cac ttg ga Arg His Leu As	t aga aaa gca gag o Arg Lys Ala Glu 85	ctg att gtg caa Leu Ile Val Gln 90	gtg 289 Val
Arg Ile Leu	Leu Lys Glu Lys	a atg gag cct gtt s Met Glu Pro Val O	caa caa ttg gag Gln Gln Leu Glu 105	ttg 337 Leu
att cat gac Ile His Asp 110	ttg aaa tat ttg Leu Lys Tyr Le 115	g ggg ctc tcg gat u Gly Leu Ser Asp 120	ttt ttt caa gat Phe Phe Gln Asp	gag 385 Glu 125
att aag gag Ile Lys Glu	atc tta ggt gt Ile Leu Gly Va 130	t ata tac aat gag l Ile Tyr Asn Glu 135	cac aaa tgc ttt His Lys Cys Phe 140	cac 433 His
aat aat gaa Asn Asn Glu	gta gag aaa at Val Glu Lys Me 145	g gat ttg tat ttc t Asp Leu Tyr Phe 150	aca gct ctt gga Thr Ala Leu Gly 155	ttc 481 Phe

aga Arg	ctc Leu	ctc Leu 160	aga Arg	caa Gln	cat His	ggt Gly	ttt Phe 165	aat Asn	att ïle	tcc Ser	caa Gln	gat Asp 170	gta Val	ttt Phe	aat Asn	529
		aag Lys														577
		aag Lys														625
		gaa Glu														673
		aaa Lys												Asn		721
		tgg Trp 240														769
caa Gln	agt Ser 255	gta Val	gag Glu	gca Ala	aga Arg	tgg Trp 260	ttc Phe	ata Ile	gat Asp	gct Ala	tat Tyr 265	gcg Ala	aga Arg	agg Arg	cca Pro	817
		aat Asn														865
		gca Ala					gaa	cta	222	_						913
		nia	Thr	His 290	Gln	Gln	Ğlu									913
		tta Leu	tgc	290 ttc	cct	gaa	Glu	Leu	Lys 295 cca	Asp	Leu gtg	Ser	Arg	Trp 300 agg	Trp	961
Ser	Arg	tta	tgc Cys 305	290 ttc Phe	cct Pro	gaa Glu gcg	Glu aag Lys gtt	ctc Leu 310	Lys 295 cca Pro	Asp ttt Phe ttt	Leu gtg Val gag	agg Arg	gat Asp 315	Trp 300 agg Arg	Trp ctc Leu cat	
Ser gtt Val	Arg gaa Glu tat	tta Leu tcc Ser	tgc Cys 305 ttc Phe	290 ttc Phe ttt Phe	cct Pro tgg Trp	gaa Glu gcg Ala	Glu aag Lys gtt Val 325 gcc	ctc Leu 310 ggg Gly	Lys 295 cca Pro atg Met	Asp ttt Phe ttt Phe	Leu gtg Val gag Glu gtt	ser agg Arg cca Pro 330	gat Asp 315 cat His	Trp 300 agg Arg caa Gln	Trp ctc Leu cat His	961
Ser gtt Val gga Gly	gaa Glu tat Tyr 335	tta Leu tcc Ser 320	tgc Cys 305 ttc Phe aga Arg	290 ttc Phe ttt Phe aaa Lys tac	cct Pro tgg Trp atg Met	gaa Glu gcg Ala gcc Ala 340	Glu aag Lys gtt Val 325 gcc Ala	ctc Leu 310 ggg Gly aca Thr	Lys 295 cca Pro atg Met att Ile	Asp ttt Phe ttt Phe att Ile	Leu gtg Val gag Glu gtt Val 345	ser agg Arg cca Pro 330 tta Leu gaa	gat Asp 315 cat His gca Ala	Trp 300 agg Arg caa Gln aca Thr	Trp  ctc Leu  cat His  gtt Val	961 1009

Pne	Thr	Asp	Thr	Phe 370	Lys	Arg	Trp	Asp	Thr 375	Glu	Ser	Ile	Thr	Arg 380	Leu	
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		gca Ala 400														1249
		cgg Arg														1297
		tgg Trp														1345
		gcc Ala														1393
		aca Thr														1441
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Leu	Tyr	Gln	Tyr	His	999 999	Ile	Leu 485 tca	Cys	Leu	Ala	Gly	Ile 490 gcg	Ile aga	Leu	Arg	1489 1537
ctt Leu gtg	Tyr ccc pro 495 ccg	Gln 480 gac	Tyr gat Asp	His ctt Leu atc	Asp ggg Gly cag	aca Thr 500	Leu 485 tca Ser	Cys tat Tyr	ttt Phe	Ala gag Glu gaa	Gly ctg Leu 505 aca	Ile 490 gcg Ala aat	Ile aga Arg	ggc Gly	gac Asp	
ctt Leu gtg Val 510	Tyr ccc Pro 495 ccg Pro	Gln 480 gac Asp	Tyr gat Asp aca Thr	His ctt Leu atc Ile	Asp 999 Gly cag Gln 515	aca Thr 500 tgc Cys	Leu 485 tca Ser tac Tyr	Cys tat Tyr atg Met	ttt Phe aag Lys	Ala gag Glu gaa Glu 520 ata	ctg Leu 505 aca Thr	Ile 490 gcg Ala aat Asn	aga Arg gct Ala	ggc Gly agt ser	gac Asp gag Glu 525	1537
ctt Leu gtg Val 510 gag Glu	Tyr ccc Pro 495 ccg Pro gag Glu	Gln 480 gac Asp aaa Lys	gat Asp aca Thr gtg Val	His ctt Leu atc Ile gag Glu 530 gcc	Asp ggg Gly cag Gln 515 cac His	aca Thr 500 tgc Cys gtg Val	Leu 485 tca Ser tac Tyr aag Lys	Cys tat Tyr atg Met ttt Phe	Leu ttt Phe aag Lys ctg Leu 535	Ala gag Glu gaa Glu 520 ata Ile	ctg Leu 505 aca Thr agg Arg	Ile 490 gcg Ala aat Asn gag Glu	aga Arg gct Ala gcg Ala	ggc Gly agt ser tgg Trp 540	gac Asp gag Glu 525 aag Lys	1537 1585
ctt Leu gtg Val 510 gag Glu gat Asp	Tyr ccc Pro 495 ccg Pro gag Glu atg Met	Gln 480 gac Asp aaa Lys gcg Ala	Tyr gat Asp aca Thr gtg Val acg Thr 545	His ctt Leu atc Ile gag Glu 530 gcc Ala gct	Asp 999 Gly cag Gln 515 cac His ata Ile	Ile aca Thr 500 tgc Cys gtg Val gca Ala att	Leu 485 tca Ser tac Tyr aag Lys gcc Ala	Cys tat Tyr atg Met ttt Phe ggt Gly 550 cgc	Leu ttt Phe aag Lys ctg Leu 535 tat Tyr	Ala gag Glu gaa Glu 520 ata Ile ccg Pro	ctg Leu 505 aca Thr agg Arg	Ile 490 gcg Ala aat Asn gag Glu ccg Pro	aga Arg gct Ala gcg Ala gat Asp 555	ggc Gly agt Ser tgg Trp 540 ggt Gly tat	gac Asp gag Glu 525 aag Lys atg Met	1537 1585 1633

gcc ggc cta ctg ttc gag cct tat gca tga acaaatggga gactgcttga

Ala Gly Leu Leu Phe Glu Pro Tyr Ala \*
590 595

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<213> Salvia officinalis

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Thr His Gln Glu Leu Lys Asp Leu Ser Arg Trp Trp Ser Arg Leu
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Cys Phe Pro Glu Lys Leu Pro Phe Val Arg Asp Arg Leu Val Glu Ser
                    310
                                         315
Phe Phe Trp Ala Val Gly Met Phe Glu Pro His Gln His Gly Tyr Gln
                                     330
Arg Lys Met Ala Ala Thr Ile Ile Val Leu Ala Thr Val Ile Asp Asp
                                 345
Ile Tyr Asp Val Tyr Gly Thr Leu Asp Glu Leu Glu Leu Phe Thr Asp
                             360
Thr Phe Lys Arg Trp Asp Thr Glu Ser Ile Thr Arg Leu Pro Tyr Tyr
                        375
                                             380
Met Gln Leu Cys Tyr Trp Gly Val His Asn Tyr Ile Ser Asp Ala Ala
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                                         395
Tyr Asp Ile Leu Lys Glu His Gly Phe Phe Cys Leu Gln Tyr Leu Arg
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                                     410
Lys Ser Val Val Asp Leu Val Glu Ala Tyr Phe His Glu Ala Lys Trp
                                 425
Tyr His Ser Gly Tyr Thr Pro Ser Leu Asp Glu Tyr Leu Asn Ile Ala
                             440
Lys Ile Ser Val Ala Ser Pro Ala Ile Ile Ser Pro Thr Tyr Phe Thr
                        455
                                             460
Phe Ala Asn Ala Ser His Asp Thr Ala Val Ile Asp Ser Leu Tyr Gln
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                                         475
Tyr His Asp Ile Leu Cys Leu Ala Gly Ile Ile Leu Arg Leu Pro Asp
                485
                                     490
Asp Leu Gly Thr Ser Tyr Phe Glu Leu Ala Arg Gly Asp Val Pro Lys
                                 505
                                                     510
Thr Ile Gln Cys Tyr Met Lys Glu Thr Asn Ala Ser Glu Glu Glu Ala
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Val Glu His Val Lys Phe Leu Ile Arg Glu Ala Trp Lys Asp Met Asn
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Thr Ala Ile Ala Ala Gly Tyr Pro Phe Pro Asp Gly Met Val Ala Gly
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ttt Pine 30	tct Ser	aac Asn	ttt Pne	tct Ser	ctt Leu 35	gac Asp	gat Asp	aag Lys	gaa Glu	caa Gln 40	caa Gln	aag Lys	tgc Cys	tca Ser	gaa Glu 45		205
acc Thr	atc Ile	gaa Glu	gca Ala	ctt Leu 50	aag Lys	caa Gln	gaa Glu	gca Ala	aga Arg 55	GTÅ GGC	atg Met	ctt Leu	atg Met	gct Ala 60	gca Ala		253
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			gca Ala													•	397
cgt Arg 110	ttc Phe	cgt Arg	ttg Leu	ctc Leu	aga Arg 115	caa Gln	cat His	caa Gln	cgc Arg	cac His 120	gtt Val	tct Ser	tgt Cys	gat Asp	gtt Val 125		445
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			gaa Glu 145														541
ttt Phe	cgc Arg	gaa Glu 160	g <b>aa</b> Glu	aga Arg	ata Ile	tta Leu	caa Gln 165	gag Glu	gct Ala	gta Val	aat Asn	ttt Phe 170	acg Thr	agg Arg	cat His		589
			gga Gly														637
			cga Arg														685
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aat Asn	ttg Leu	tat Tyr 240	aag Lys	gaa Glu	gag Glu	ctc Leu	tcc Ser 245	caa Gln	ctc Leu	tcc Ser	agg Arg	tgg Trp 250	tgg Trp	aac Asn	aca Thr	329
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gat Asp																1021
caa Gln	gtc Val	tta Leu 320	gac Asp	aag Lys	tgg Trp	gat Asp	aga Arg 325	gat Asp	gaa Glu	gct Ala	gaa Glu	cga Arg 330	ctc Leu	cca Pro	gaa Glu	1069
Gln	Val atg	Leu 320 aaa	Asp	Lys	Trp	gat Asp cga Arg 340	Arg 325 ttt	Asp	Glu	Ala	Glu	Arg 330 tat	Leu gaa	Pro	Glu	1069
Gln tac Tyr	Val atg Met 335	Leu 320 aaa Lys gat	Asp atc Ile	Lys gtt Val gcg	Trp tat Tyr	Asp cga Arg 340 ctt	Arg 325 ttt Phe	Asp att Ile aaa	Glu ttg Leu agc	Ala agt Ser	Glu ata Ile 345 gca	Arg 330 tat Tyr	Leu gaa Glu cct	Pro aat Asn tat	Glu tat Tyr	
Gln tac Tyr gaa Glu 350	Val atg Met 335 cgt Arg	Leu 320 aaa Lys gat Asp	Asp atc Ile gca Ala	Lys gtt Val gcg Ala aaa	Trp tat Tyr aaa Lys 355 caa	Asp cga Arg 340 ctt	Arg 325 ttt Phe gga Gly	Asp att Ile aaa Lys	Glu ttg Leu agc Ser	Ala agt ser ttt Phe 360	Glu ata Ile 345 gca Ala	Arg 330 tat Tyr gct Ala	gaa Glu cct Pro	Pro aat Asn tat Tyr	Glu tat Tyr ttt Phe 365 aag	1117
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					aaa Lys											1501
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 Lys
 His
 Ala
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 Trp
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 Phe
 Ser
 Asn

 Phe
 Ser
 Leu
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 Asp
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 Glu
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 Cys
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 Arg
 Gly
 Met
 Leu
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 Ala
 Ala
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 Thr
 Thr
 Thr
 Pro

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Leu Gln Gln Met Thr Leu Ile Asp Thr Leu Glu Arg Leu Gly Leu Ser Phe His Phe Glu Thr Glu Ile Glu Tyr Lys Ile Glu Leu Ile Asn Ala Ala Glu Asp Asp Gly Phe Asp Leu Phe Ala Thr Ala Leu Arg Phe Arg Leu Leu Arg Gln His Gln Arg His Val Ser Cys Asp Val Phe Asp Lys 1.20 Phe Ile Asp Lys Asp Gly Lys Phe Glu Glu Ser Leu Ser Asn Asn Val Glu Gly Leu Leu Ser Leu Tyr Glu Ala Ala His Val Gly Phe Arg Glu Glu Arg Ile Leu Gln Glu Ala Val Asn Phe Thr Arg His His Leu Glu Gly Ala Glu Leu Asp Gln Ser Pro Leu Leu Ile Arg Glu Lys Val Lys Arg Ala Leu Glu His Pro Leu His Arg Asp Phe Pro Ile Val Tyr Ala Arg Leu Phe Ile Ser Ile Tyr Glu Lys Asp Asp Ser Arg Asp Glu Leu Leu Leu Lys Leu Ser Lys Val Asn Phe Lys Phe Met Gln Asn Leu Tyr Lys Glu Glu Leu Ser Gln Leu Ser Arg Trp Trp Asn Thr Trp Asn Leu Lys Ser Lys Leu Pro Tyr Ala Arg Asp Arg Val Val Glu Ala Tyr Val Trp Gly Val Gly Tyr His Tyr Glu Pro Gln Tyr Ser Tyr Val Arg Met Gly Leu Ala Lys Gly Val Leu Ile Cys Gly Ile Met Asp Asp Thr Tyr Asp Asn Tyr Ala Thr Leu Asn Glu Ala Gln Leu Phe Thr Gln Val Leu Asp Lys Trp Asp Arg Asp Glu Ala Glu Arg Leu Pro Glu Tyr Met Lys Ile Val Tyr Arg Phe Ile Leu Ser Ile Tyr Glu Asn Tyr Glu Arg Asp Ala Ala Lys Leu Gly Lys Ser Phe Ala Ala Pro Tyr Phe Lys Glu Thr Val Lys Gln Leu Ala Arg Ala Phe Asn Glu Glu Gln Lys Trp Val Met Glu Arg Gln Leu Pro Ser Phe Gln Asp Tyr Val Lys Asn Ser Glu Lys Thr Ser Cys Ile Tyr Thr Met Phe Ala Ser Ile Ile Pro Gly Leu Lys Ser Val Thr Gln Glu Thr Ile Asp Trp Ile Lys Ser Glu Pro Thr Leu Ala Thr Ser Thr Ala Met Ile Gly Arg Tyr Trp Asn Asp Thr Ser Ser Gln Leu Arg Glu Ser Lys Gly Gly Glu Met Leu Thr Ala Leu Asp Phe His Met Lys Glu Tyr Gly Leu Thr Lys Glu Glu Ala Ala Ser Lys Phe Glu Gly Leu Val Glu Glu Thr Trp Lys Asp Ile Asn Lys Glu Phe Ile

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			500					505					510	Phe	Leu		
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Val 545	Asp	Ala	Ile	Val	Phe 550						3.0						
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aga Arg	aca Thr	atc Ile	cca Pro	aat Asn 35	ctt Leu	gga Gly	atg Met	cgt Arg	agg Arg 40	cga Arg	ggg Gly	a <b>aa</b> Lys	tct Ser	gtc Val 45	acg Thr		206
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Pro 335	aaa Lys	ttg Leu	aca Thr	ttt Phe	gct Ala 340	cgg Arg	cat His	cgt Arg	cat His	gtg Val 345	gaa Glu	ttc Phe	tac Tyr	act Thr	ttg Leu 350		1113
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	gcc Ala																1214
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	gct Ala																1454
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Leu Arg Ser Asn Asp Asn Ile Pro Met Leu Ala Lys Lys His Ala Phe
575
                                         585
gac ata aca aga get etc cac cat etc tac ata tat ega gat gge ttt
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Glu Ser Met Leu Phe
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atcgtgtggg tatgattaaa aagctagagc ttactaggtt agtaacatgg tgataaaagt
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			Cys.	165					170					175	Thr
			Phe 180					185					190		
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	210		Thr			215					220			_	_
225			Ile		230					235					240
			Thr	245					250			_		255	
			Ser 260					265				_	270	-	
		275	Pro				280					285			
	290		Ser			295					300	_	_		
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	N.		Tyr	325					330					335	-
			Ala 340					345			_		350		
cya	ire	355	Ile	Asp	Pro	гЛа	360	ser	Ala	Pne	Arg	Leu 365	GIY	Phe	Ala
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Gly 385	370 Thr	Ile	Asp	Glu	Leu 390	375 Glu	Val Leu	Phe	Thr	Ser	380 Ala	Tyr Ile	Lys	Arg	Trp
Gly 385 Asn	370 Thr Ser	Ile Ser	Asp Glu	Glu Ile 405	Leu 390 Glu	375 Glu His	Val Leu Leu	Phe Pro	Thr Glu 410	Ser 395 Tyr	380 Ala Met	Tyr Ile Lys	Lys	Arg Val 415	Trp 400 Tyr
Gly 385 Asn Met	370 Thr Ser Val	Ile Ser Val	Asp Glu Phe 420	Glu Ile 405 Glu	Leu 390 Glu Thr	375 Glu His Val	Val Leu Leu Asn	Phe Pro Glu 425	Thr Glu 410 Leu	Ser 395 Tyr Thr	380 Ala Met Arg	Tyr Ile Lys Glu	Lys Cys Ala 430	Arg Val 415 Glu	Trp 400 Tyr Lys
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acc Thr	cca Pro	gct Ala	gct Ala 5	gta Val	gta Val	atg Met	agt Ser	aac Asn 10	tac Tyr	gga Gly	gag Glu	gag Glu	gag Glu 15	att Ile	gtt Val	107
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			ctt Leu													539
			gta Val 165													587
			act Thr													635
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			aga Arg													731,
			aag Lys													779
			cag Gln 245													827
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	gta Val																1595
	act Thr 515																1643
	aat Asn																1691

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Leu Tyr Asn Asp Tyr Glu Met Glu Leu Ser Lys Asp Gly Arg Ser Asp
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Val Val His Tyr Ala Lys Glu Arg Met Lys Glu Ile Val Arg Asn Tyr
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Trp Leu Ala Lys Asn Pro Lys Ile Leu Glu Ala Asn Val Thr Leu Cys
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                                                 445
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Gln Ile Ala Thr Gly Ile Glu Cys Tyr Met Arg Asp Tyr Gly Val Ser
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Thr Glu Lys Ala Met Glu Lys Phe Gln Glu Met Ala Glu Thr Ala Trp
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Lys Asp Val Asn Glu Gly Ile Leu Arg Pro Thr Pro Val Ser Thr Glu
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Ile Leu Thr Arg Ile Leu Asn Leu Ala Arg Ile Ile Asp Val Thr Tyr
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Glu Leu Lys Ala Gln Val Arg Lys Met Ile M 55 60	aty gaa cct gtt gat gat 1750 Met Glu Pro Val Asp Asp 65
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cgt gac acc aac aac aat gat gcg gac acc g Arg Asp Thr Asn Asn Asn Asp Ala Asp Thr A 105	
ctt cga ttc cgg tta ctt aga gag cat ggc t Leu Arg Phe Arg Leu Leu Arg Glu His Gly F 120 125	
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Asp Glu Ala Gly Asn Phe Lys Ala Ser Leu 140 145  ttg ttg gaa ctt tat gaa gct tcc tat atg a Leu Leu Glu Leu Tyr Glu Ala Ser Tyr Met A	Thr Ser Asp Val Gln Gly 150 agg gtc cat ggg gaa gat 2208
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ccc caa tac tct ctt ggt aga aag atg ttg aca aaa gtc ata gca atg Pro Gln Tyr Ser Leu Gly Arg Lys Met Leu Thr Lys Val Ile Ala Met 290 295 300	2699
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Arg Trp Asp 330	2971
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att aaa tgc atg aac caa ctc ccg aat tac atg aaa ata agc tac aag Ile Lys Cys Met Asn Gln Leu Pro Asn Tyr Met Lys Ile Ser Tyr Lys 335  gca cta tta gat gtt tat gaa gaa atg gaa cag ctg ttg gca aat caa Ala Leu Leu Asp Val Tyr Glu Glu Met Glu Gln Leu Leu Ala Asn Gln 350  ggg aga cag tac cga gtt gag tat gcg aaa aag gcg gtatgtaatg Gly Arg Gln Tyr Arg Val Glu Tyr Ala Lys Lys Ala	<ul><li>2971</li><li>3019</li><li>3067</li></ul>

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gct ata acg gcg ttt gtc ggc atg ggc gaa gtt ata acc cct gag acc Ala Ile Thr Ala Phe Val Gly Met Gly Glu Val Ile Thr Pro Glu Thr 415 420 425 430	3370
ttc aaa tgg gcc gcc agt gac ccc aag atc att aag gct tcc acc att Phe Lys Trp Ala Ala Ser Asp Pro Lys Ile Ile Lys Ala Ser Thr Ile 435 440 445	3418
att tgc agg ttc atg gac gat att gct gaa cat aag gtatactata Ile Cys Arg Phe Met Asp Asp Ile Ala Glu His Lys 450 455	3464
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<211> 47

<212> PRT

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<400> 39

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 Leu
 Val
 Gln
 Ala
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 Glu
 Ala
 Lys
 Trp
 Thr
 His

 Gln
 Asn
 Tyr
 Lys
 Pro
 Thr
 Phe
 Glu
 Glu
 Phe
 Arg
 Asn
 Ala
 Leu
 Pro

 Thr
 Ser
 Gly
 Tyr
 Ala
 Met
 Leu
 Ala
 Ile
 Thr
 Ala
 Phe
 Val
 Gly
 Met
 Gly

 Glu
 Val
 Ile
 Thr
 Phe
 Lys
 Trp
 Ala
 Ser
 Asp
 Pro
 Lys

 Follow
 Tyr
 Ala
 Ser
 Thr
 Ile
 Ile
 Cys
 Arg
 Phe
 Met
 Asp
 Ile
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<211> 97

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Ile															

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<213> Ricinus communis

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		ttc Phe																540
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		gaa Glu															-	684
		tcc Ser															•	732
		ttc Phe																780
		ctc Leu 240																828
		gtg Val																876
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		tgg Trp												Tyr				1020

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g <b>ag</b> Glu	aag Lys	ctg Leu 385	cca Pro	gat Asp	tac Tyr	atg Met	aaa Lys 390	gtt Val	att Ile	tat Tyr	aaa Lys	cta Leu 395	ttg Leu	cta Leu	aac Asn		1260
acc Thr	ttc Phe 400	tct Ser	gaa Glu	ttc Phe	gag Glu	aaa Lys 405	gaa Glu	ttg Leu	acg Thr	gcg Ala	gaa Glu 410	ggc Gly	aag Lys	tcc Ser	tac Tyr	-	1308
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tac Tyr	ctg Leu	g <b>ag</b> Glu	gct Ala	gta Val 435	tgg Trp	cgc Arg	gac Asp	gag Glu	ggt Gly 440	aaa Lys	ata Ile	cca Pro	tcg Ser	ttc Phe 445	gat Asp		1404
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<212> PRT

<213> Ricinus communis

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Ser Leu Arg Gly Asp Ala Lys Gly Met Leu Ser Leu Phe Glu Ala Ser His Leu Ser Val His Gly Glu Asp Ile Leu Glu Glu Ala Phe Ala Phe Thr Lys Asp Tyr Leu Gln Ser Ser Ala Val Glu Leu Phe Pro Asn Leu Lys Arg His Ile Thr Asn Ala Leu Glu Gln Pro Phe His Ser Gly Val Pro Arg Leu Glu Ala Arg Lys Phe Ile Asp Leu Tyr Glu Ala Asp Ile Glu Cys Arg Asn Glu Thr Leu Leu Glu Phe Ala Lys Leu Asp Tyr Asn Arg Val Gln Leu Leu His Gln Gln Glu Leu Cys Gln Phe Ser Lys Trp Trp Lys Asp Leu Asn Leu Ala Ser Asp Ile Pro Tyr Ala Arg Asp Arg Met Ala Glu Ile Phe Phe Trp Ala Val Ala Met Tyr Phe Glu Pro Asp Tyr Ala His Thr Arg Met Ile Ile Ala Lys Val Val Leu Leu Ile Ser Leu Ile Asp Asp Thr Ile Asp Ala Tyr Ala Thr Met Glu Glu Thr His Ile Leu Ala Glu Ala Val Ala Arg Trp Asp Met Ser Cys Leu Glu Lys Leu Pro Asp Tyr Met Lys Val Ile Tyr Lys Leu Leu Leu Asn Thr Phe Ser Glu Phe Glu Lys Glu Leu Thr Ala Glu Gly Lys Ser Tyr Ser Val Lys Tyr Gly Arg Glu Ala Phe Gln Glu Leu Val Arg Gly Tyr Tyr Leu Glu Ala Val Trp Arg Asp Glu Gly Lys Ile Pro Ser Phe Asp Asp Tyr Leu Tyr Asn Gly Ser Met Thr Thr Gly Leu Pro Leu Val Ser Thr Ala Ser Phe Met Gly Val Gln Glu Ile Thr Gly Leu Asn Glu Phe Gln Trp Leu Glu Thr Asn Pro Lys Leu Ser Tyr Ala Ser Gly Ala Phe Ile Arg Leu Val Asn Asp Leu Thr Ser His Val Thr Glu Gln Gln Arg Gly His Val Ala Ser Cys Ile Asp Cys Tyr Met Asn Gln His Gly Val Ser Lys Asp Glu Ala Val Lys Ile Leu Gln Lys Met Ala Thr Asp Cys Trp Lys Glu Ile Asn Glu Glu Cys Met Arg Gln Ser Gln Val Ser Val Gly His Leu Met Arg Ile Val Asn Leu Ala Arg Leu Thr Asp Val Ser Tyr Lys Tyr Gly Asp Gly Tyr Thr Asp Ser Gln Gln Leu Lys Gln Phe Val Lys Gly Leu Phe Val Asp Pro Ile Ser Ile 

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gaa Glu	gct Ala 610	gaa Glu	tgg Trp	tta Leu	gct Ala	gct Ala 615	gag Glu	tat Tyr	gtg Val	cct Pro	acc Thr 620	ttg Leu	gac Asp	gag Glu	tac Tyr		1873
ata Ile 625	aag Lys	aat Asn	gga Gly	atc Ile	aca Thr 630	tct Ser	atc Ile	ggc Gly	caa Gln	cgt Arg 635	ata Ile	ctt Leu	ctg Leu	ttg Leu	agt Ser 640		1921
gga Gly	gtg Val	ttg Leu	ata Ile	atg Met 645	gat Asp	gly aaa	caa Gln	ctc Leu	ctt Leu 650	tcg Ser	caa Gln	gag Glu	gca Ala	tta Leu 655	gag Glu		1969
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cgt gga Arg Gly 690	gaa ttg Glu Leu	gcg tcc Ala Ser	agc at Ser Ile 695	E gaa ∍ Glu	tgt Cys	tac Tyr	atg Met 700	aaa Lys	gac Asp	cat His	cct Pro		2113
gaa tgt Glu Cys 705	aca gag Thr Glu	gaa gag Glu Glu 710	gct cte Ala Le	gat 1 Asp	cac His	atc Ile 715	tat Tyr	agc Ser	att Ile	ctg Leu	gag Glu 720		2161
ccg gcg Pro Ala	gtg aag Val Lys	gaa ctg Glu Leu 725	aca aga Thr Arg	a gag g Glu	ttt Phe 730	ctg Leu	aag Lys	ccc Pro	gac Asp	gac Asp 735	gtc Val		2203
cca ttc Pro Phe	gcc tgc Ala Cys 740	aag aag Lys Lys	atg ct Met Le	ttc Phe 745	gag Glu	gag Glu	aca Thr	aga Arg	gtg Val 750	acg Thr	atg Met		2257
gtg ata Val Ile	ttc aag Phe Lys 755	gat gga Asp Gly	gat gga Asp Gly 760	y Phe	ggt Gly	gtt Val	tcc Ser	aaa Lys 765	tta Leu	gaa Glu	gtc Val		2305
aaa gat Lys Asp 770	cat atc His Ile	aaa gag Lys Glu	tgt cto Cys Let 775	att Ile	gaa Glu	ccg Pro	ctg Leu 780	cca Pro	ctg Leu	taa *		-	2350
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<213> Abies grandis

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145					150					155					160
			Tyr	165					170					175	Arg
			Leu 180					185					190	Val	
		195	Leu				200					205			-
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			Leu	245					250					255	Pro
			Pro 260					265					270		
		275	Leu				280					285		_	
	290		Tyr			295					300				
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			Arg	325					330					335	_
			Asp 340					345					350		
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_	370		Gly			375					380				
Lys 385			Arg		390					395					400
			Gln	405					410					415	
			Ala 420					425				_	430	_	
Val		435	Pro				440			_	_	445	_	-	
	450		Asn			455					460			_	
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			Asp	485					490					495	
			Phe 500					505					510		
		515	Сув				520					525			
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545			Glu		550					555					560
Leu	LLO	Asp	ıyr	met	Lys	Leu	Cvs	Tvr	Gln	Ile	Tvr	Tvr	Asp	Ile	Val

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Ser Phe Phe Arg Lys Gly Trp Glu Asp Tyr Leu Leu Gly Tyr Tyr Glu
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Glu Ala Glu Trp Leu Ala Ala Glu Tyr Val Pro Thr Leu Asp Glu Tyr
                        615
                                             620
Ile Lys Asn Gly Ile Thr Ser Ile Gly Gln Arg Ile Leu Leu Leu Ser
625
                    €30
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Gly Val Leu Ile Met Asp Gly Gln Leu Leu Ser Gln Glu Ala Leu Glu
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Lys Val Asp Tyr Pro Gly Arg Arg Val Leu Thr Glu Leu Asn Ser Leu
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Ile Ser Arg Leu Ala Asp Asp Thr Lys Thr Tyr Lys Ala Glu Lys Ala
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Arg Gly Glu Leu Ala Ser Ser Ile Glu Cys Tyr Met Lys Asp His Pro
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Pro Phe Ala Cys Lys Lys Met Leu Phe Glu Glu Thr Arg Val Thr Met
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cac gga aat gtg tgg gac gat gac ctc ata cac tct ctc aac tcg ccc
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His Gly Asn Val Trp Asp Asp Leu Ile His Ser Leu Asn Ser Pro
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tat ggg gca cct gca tat tat gag ctc ctt caa aag ctt att cag gag
                                                                       144
Tyr Gly Ala Pro Ala Tyr Tyr Glu Leu Leu Gln Lys Leu Ile Gln Glu
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atc aag cat tta ctt ttg act gaa atg gaa atg gat gat ggc gat cat
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Ile Lys His Leu Leu Leu Thr Glu Met Glu Met Asp Asp Gly Asp His
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atc Ile	gat Asp	aga Arg	cat His	ttt Phe 85	gaa Glu	cac His	gaa Glu	ata Ile	caa Gln 90	aca Thr	gct Ala	gct Ala	tta Leu	gat Asp 95	tac Tyr		288
gtt Val	tac Tyr	aga Arg	tgg Trp 100	tgg Trp	aac Asn	gaa Glu	aaa Lys	ggt Gly 105	atc Ile	gly aaa	gag Glu	gga Gly	tca Ser 110	aga Arg	gat Asp		336
tcc Ser	ttc Phe	agc Ser 115	aaa Lys	gat Asp	ctg Leu	aac Asn	gct Ala 120	acg Thr	gct Ala	tta Leu	gga Gly	ttt Phe 125	cgc Arg	gct Ala	ctc Leu		384
cga Arg	ctg Leu 130	cat His	cga Arg	tat Tyr	aac Asn	gta Val 135	tcg Ser	tca Ser	ggt Gly	gtg Val	ttg Leu 140	aag Lys	aat Asn	ttc Phe	aag Lys		432
gat Asp 145	gaa Glu	aac Asn	gly aga	aag Lys	ttc Phe 150	ttc Phe	tgc Cys	aac Asn	ttt Phe	act Thr 155	ggt Gly	gaa Glu	gaa Glu	gga Gly	aga Arg 160		480
gga Gly	gat Asp	aaa Lys	caa Gln	gtg Val 165	aga Arg	agc Ser	atg Met	ttg Leu	tcg Ser 170	tta Leu	ctt Leu	cga Arg	gct Ala	tca Ser 175	gag Glu	•	528
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gtg Val	gat Asp 210	caa Gln	agc Ser	ctt Leu	ttg Leu	aga Arg 215	gag Glu	gtg Val	aag Lys	tac Tyr	gca Ala 220	ttg Leu	gag Glu	ttt Phe	cca Pro		672
tgg Trp 225	cat His	tgc Cys	agt Ser	gtg Val	ccg Pro 230	aga Arg	tgg Trp	gag Glu	gca Ala	agg Arg 235	agc Ser	ttt Phe	ctc Leu	gaa Glu	ata Ile 240		720
	gga Gly																768
	aag Lys																816

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cag Gln	ctg Leu 290	aat Asn	ttc Phe	tat Tyr	cga Arg	aag Lys 295	cga Arg	cac His	gtg Val	gaa Glu	tat Tyr 300	tat Tyr	tct Ser	tgg Trp	gtt Val		912
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cat His	att Ile	ccc Pro	act Thr 420	ttt Phe	aac Asn	gaa Glu	tac Tyr	ata Ile 425	aag Lys	aac Asn	ggc	atg Met	gct Ala 430	agc Ser	tca Ser		1296
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gac Asp 465	ctc Leu	tta Leu	gaa Glu	ttg Leu	acg Thr 470	ggc Gly	aga Arg	atc Ile	gcc Ala	gat Asp 475	gac Asp	tta Leu	aaa Lys	gat Asp	ttc Phe 480		1440
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tct Ser	gac Asp	aag Lys	gaa Glu	gta Val 565	aag Lys	gac Asp	cag Gln	ata Ile	ttc Phe 570	aaa Lys	att Ile	cta Leu	gtc Val	cac His 575	caa Gln	1728
gtt Val	cca Pro	atg Met	gag Glu 580	gaa Glu	tag *	tgat	ggto	ett g	gttg	gtagt	t gt	ctat	tato	J		1776
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<213> Abies grandis

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Arg Leu His Arg Tyr Asn Val Ser Ser Gly Val Leu Lys Asn Phe Lys

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Giy	Asp	rys	GIII	165	Arg	261	MEC	Leu	170	Leu	Leu	Arg	Ala	175	GII
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Val	Asp 210	Gln	Ser	Leu	Leu	Arg 215	Glu	Val	Lys	Tyr	Ala 220	Leu	Glu	Phe	Pro
Trp 225	His	Cya	Ser	Val	Pro 230	Arg	Trp	Glu	Ala	Arg 235	Ser	Phe	Leu	Glu	Ile 240
Tyr	Gly	His	Asn	His 245	Ser	Trp	Leu	TAa	Ser 250	Asn	Ile	Asn	Gln	Lys 255	Met
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Gln	Leu 290	Asn	Phe	Tyr	Arg	Lys 295	Arg	His	Val	Glu	Tyr 300	Tyr	Ser	Trp	Val
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Lvs	Ara	$\alpha$	Clar	Ara	Asp	Met	Thr	Thr	Ile	Val	Lys	Asp	Cys	Trp	Lys
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385 Arg	Tyr	Ile	Glu	Ser 405	Tyr	Leu	Gln	Glu	Ala 410	395 Glu	_	Ile	Ala	Thr 415	400 Gly
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Asp 465 Glu Met	Tyr Ile Met Pro 450 Leu Asp	Ile Pro Cys 435 Asp Leu Glu	Glu Thr 420 Ile Asn Glu Lys Asn 500	Ser 405 Phe Leu Ile Leu Glu 485 Pro	Tyr Asn Asn Leu Thr 470 Arg Glu	Leu Glu Leu Glu 455 Gly Gly Ser	Gln Tyr Asn 440 Gln Arg Glu Thr	Glu Ile 425 Pro Ile Ile Met Val 505	Ala 410 Lys Leu His Ala Ala 490 Glu	395 Glu Asn Leu Ser Asp 475 Ser Asn	Gly Leu Pro 460 Asp Ser	Ile Met Leu 445 Ser Leu Leu Leu	Ala Ala 430 Asp Lys Lys Gln Asn 510	Thr 415 Ser Lys Ile Asp Cys 495 His	Gly Ser Leu Leu Phe 480 Tyr Ile
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Asp Asp Arg Ile Gln Ser Leu Asn Ser Pro Tyr Gly Ala Pro Ala Tyr
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Gln Glu Arg Ser Glu Lys Leu Ile Glu Glu Ile Lys Leu Phe Leu
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Ser Asp Met Asp Asp Ser Cys Asn Asp Ser Asp Arg Asp Leu Ile Lys
cgt ctt gag atc gtt gat act gtc gag tgt ctg gga att gat cga cat
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Arg Leu Glu Ile Val Asp Thr Val Glu Cys Leu Gly Ile Asp Arg His
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Phe Gln Pro Glu Ile Lys Leu Ala Leu Asp Tyr Val Tyr Arg Cys Trp
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Asn Glu Arg Gly Ile Gly Glu Gly Ser Arg Asp Ser Leu Lys Lys Asp
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agt Ser 240	gtg Val	cag Gln	aga Arg	tgg Trp	gag Glu 245	gca Ala	agg Arg	agc Ser	ttt Phe	atc Ile 250	gaa Glu	ata Ile	ttt Phe	gga Gly	caa Gln 255	768
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gcg Ala	aaa Lys	ttg Leu	gac Asp 275	ttc Phe	aat Asn	att Ile	ctg Leu	caa Gln 280	tgc Cys	aca Thr	cat His	cag Gln	aaa Lys 285	gaa Glu	ctg Leu	864
cag Gln	att Ile	atc Ile 290	tca Ser	agg Arg	tgg Trp	ttc Phe	gca Ala 295	gac Asp	tca Ser	agt Ser	ata Ile	gca Ala 300	tcc Ser	ctg Leu	aat Asn	912
ttc Phe	tat Tyr 305	egg Arg	aaa Lys	tgt Cys	tac Tyr	gtc Val 310	gaa Glu	ttt Phe	tac Tyr	ttt Phe	tgg Trp 315	atg Met	gct Ala	gca Ala	gcc Ala	960
atc Ile 320	tcc Ser	gag Glu	ccg Pro	gag Glu	ttt Phe 325	tct Ser	gga Gly	agc Ser	aga Arg	gtt Val 330	gcc Ala	ttc Phe	aca Thr	aaa Lys	att Ile 335	1008
gct Ala	ata Ile	ctg Leu	atg Met	aca Thr 340	atg Met	cta Leu	gat Asp	gac Asp	ctg Leu 345	tac Tyr	gat Asp	act Thr	cac	gga Gly 350	acc Thr	1056
ttg Leu	gac Asp	caa Gln	ctc Leu	aaa Lys	atc Ile	ttt Phe	aca Thr	gag Glu	gga Gly	gtg Val	aga Arg	cga Arg	tgg Trp	gat Asp	gtt Val	1104

tcg ttg g Ser Leu V	gta gag Val Glu 1870	ggc ctc Gly Leu	Pro A	gac Asp 375	ttc Phe	atg Met	aaa Lys	att Ile	gca Ala 380	ttc Phe	gag Glu	ttc Phe		1152
tgg tta a Trp Leu I 385														1200
ggg caa g Gly Gln A 400														1248
ctt gaa g Leu Glu A	gct tat Ala Tyr	ctg caa Leu Gln 420	gat o	gcg Ala	gaa Glu	tgg Trp 425	ata Ile	gcc Ala	act Thr	gga Gly	cat His 430	gtc Val		1296
ccc acc t Pro Thr F				Asn .										1344
tgt gta t Cys Val I			Pro											1392
atc gac a Ile Asp I 465													•	1440
att gaa t Ile Glu I 480	tg gct Leu Ala	tcc agg Ser Arg 485	ctc ( Leu	gtc Val	gat Asp	gac Asp	gcg Ala 490	aga Arg	gat Asp	ttc Phe	cag Gln	gcg Ala 495	-	1488
gag aag g Glu Lys A														1536
cat cct o				Asp										1584
ctt ggc a Leu Gly A			Glu											1632
gac agt o Asp Ser V 545	gtg cca Val Pro	ctc tcg Leu Ser	tgt Cys 550	aag Lys	aag Lys	tac Tyr	agc Ser	ttc Phe 555	cat His	gta Val	ttg Leu	gca Ala		1680
cga agc a Arg Ser 1 560	atc caa Ile Gln	ttc atg Phe Met 565	Tyr .	aat Asn	caa Gln	ggc Gly	gat Asp 570	ggc Gly	ttc Phe	tcc Ser	att Ile	tcg Ser 575		1728

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aac aaa gtg atc aag gat caa gtg cag aaa gtt ctt att gtc ccc gtg
Asn Lys Val Ile Lys Asp Gln Val Gln Lys Val Leu Ile Val Pro Val
580 585 590

cct att tga
Pro Ile *
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290
                        295
                                             300
Tyr Arg Lys Cys Tyr Val Glu Phe Tyr Phe Trp Met Ala Ala Ile
                    310
                                         315
Ser Glu Pro Glu Phe Ser Gly Ser Arg Val Ala Phe Thr Lys Ile Ala
                325
                                     330
Ile Leu Met Thr Met Leu Asp Asp Leu Tyr Asp Thr His Gly Thr Leu
            340
                                 345
                                                     350
Asp Gln Leu Lys Ile Fhe Thr Glu Gly Val Arg Arg Trp Asp Val Ser
                             360
                                                 365
Leu Val Glu Gly Leu Pro Asp Phe Met Lys Ile Ala Phe Glu Phe Trp
                        375
Leu Lys Thr Ser Asn Glu Leu Ile Ala Glu Ala Val Lys Ala Gln Gly
                    390
                                         395
Gln Asp Met Ala Ala Tyr Ile Arg Lys Asn Ala Trp Glu Arg Tyr Leu
                405
                                     410
                                                         415
Glu Ala Tyr Leu Gln Asp Ala Glu Trp Ile Ala Thr Gly His Val Pro
                                 425
            420
Thr Phe Asp Glu Tyr Leu Asn Asn Gly Thr Pro Asn Thr Gly Met Cys
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                             440
Val Leu Asn Leu Ile Pro Leu Leu Leu Met Gly Glu His Leu Pro Ile
                        455
                                             460
Asp Ile Leu Glu Gln Ile Phe Leu Pro Ser Arg Phe His His Leu Ile
                                         475
                    470
Glu Leu Ala Ser Arg Leu Val Asp Asp Ala Arg Asp Phe Gln Ala Glu
                485
                                     490
Lys Asp His Gly Asp Leu Ser Cys Ile Glu Cys Tyr Leu Lys Asp His
            500
                                 505
                                                     510
Pro Glu Ser Thr Val Glu Asp Ala Leu Asn His Val Asn Gly Leu Leu
                             520
                                                 525
Gly Asn Cys Leu Leu Glu Met Asn Trp Lys Phe Leu Lys Lys Gln Asp
                        535
                                             540
Ser Val Pro Leu Ser Cys Lys Lys Tyr Ser Phe His Val Leu Ala Arg
                    550
                                         555
Ser Ile Gln Phe Met Tyr Asn Gln Gly Asp Gly Phe Ser Ile Ser Asn
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<sup>&</sup>lt;211> 2024

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Lycopersicon esculentum

<sup>&</sup>lt;220>

<sup>&</sup>lt;221> CDS -

<sup>&</sup>lt;222> (32)...(1678)

<sup>&</sup>lt;223> VFNT germacrene C synthase

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ttc Phe	ctt Leu 25	tct Ser	tat Tyr	act Thr	cat His	gaa Glu 30	att Ile	act Thr	aat Asn	caa Gln	gaa Glu 35	aaa Lys	gtt Val	gaa Glu	gtt Val		148
gat Asp 40	gag Glu	tac Tyr	aaa Lys	gag Glu	aca Thr 45	att Ile	aga Arg	aaa Lys	atg Met	ctg Leu 50	gtg Val	gaa Glu	act Thr	tgc Cys	gac Asp 55		196
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gtg Val	gct Ala	tat Tyr	cat His 75	ttc Phe	gat Asp	aat Asn	gaa Glu	att Ile 80	gaa Glu	aca Thr	tcc Ser	att Ile	caa Gln 85	aac Asn	att Ile		292
ttt Phe	gat Asp	gca Ala 90	tcg Ser	tcc Ser	aaa Lys	cag Gln	aat Asn 95	gat Asp	aat Asn	gac Asp	aac Asn	aac Asn 100	ctt Leu	tac Tyr	gtt Val	٠	340
gtg Val	tct Ser 105	ctt Leu	cgt Arg	ttt Phe	cga Arg	ctt Leu 110	gtg Val	agg Arg	caa Gln	caa Gln	ggc Gly 115	cat His	tac Tyr	atg Met	tct Ser	•	388
tca Ser 120	gat Asp	gtg Val	ttc Phe	aag Lys	caa Gln 125	ttc Phe	acc Thr	aac Asn	caa Gln	gat Asp 130	ggg Gly	aaa Lys	ttc Phe	aag Lys	gaa Glu 135		436
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cat	ctg Leu	aga Arg	gtg Val 155	cgt Arg	aat Asn	gag Glu	gag Glu	att Ile 160	ctt Leu	gaa Glu	g <b>aa</b> Glu	gct Ala	ctt Leu 165	aca Thr	ttt Phe		532
	acc Thr																580
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atg Met 200	act Thr	tta Leu	cca Pro	agg Arg	atg Met 205	gga Gly	gct Ala	aga Arg	aaa Lys	tac Tyr 210	ata Ile	tcc Ser	att Ile	tac Tyr	gaa Glu 215		676

aac a Asn A	aat Asn	gat Asp	gca Ala	cac His 220	cac His	cat His	ttg Leu	ctt Leu	ttg Leu 225	aaa Lys	ttt Phe	gct Ala	aaa Lys	ttg Leu 230	gat Asp		724
ttt a Phe A	aac Asn	atg Met	ctg Leu 235	caa Gln	aag Lys	ttt Phe	cac His	caa Gln 240	aga Arg	gag Glu	ctt Leu	agt Ser	gat Asp 245	ctt Leu	aca Thr		772
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gac a Asp A	agg Arg 265	ttg Leu	gtt Val	gag Glu	tgt Cys	tac Tyr 270	ttc Phe	tgg Trp	ata Ile	tta Leu	gga Gly 275	gtg Val	tat Tyr	ttt Phe	gag Glu		868
cca a Pro I 280																	916
acc t Thr S	Ser	att Ile	att Ile	gac Asp 300	gac Asp	act Thr	ttt Phe	gat Asp	gct Ala 305	tat Tyr	gca Ala	acc Thr	ttt Phe	gac Asp 310	g <b>aa</b> Glu	-	964
ctt g Leu V	gtg Val	act Thr	ttc Phe 315	aat Asn	gat Asp	gca Ala	atc Ile	cag Gln 320	aga Arg	tgg Trp	gat Asp	gct Ala	aat Asn 325	gca Ala	att Ile	•	1012
gat t Asp S																	1060
att t Ile 1	tac Tyr 345	agt Ser	gaa Glu	atg Met	gaa Glu	caa Gln 350	gtg Val	ttg Leu	tcc Ser	aaa Lys	gaa Glu 355	ggt Gly	aaa Lys	ctg Leu	gac Asp		1108
cgt g Arg V 360																	1156
ttt a Phe I	aag Lys	gaa Glu	acc Thr	caa Gln 380	tgg Trp	ttg Leu	aat Asn	gat Asp	tgt Cys 385	gac Asp	cat His	att Ile	cca Pro	aaa Lys 390	tat Tyr		1204
gag g Glu G																	1252
tca a																	1300

ttt gaa Phe Glu 425	tgg ttg Trp Leu	atg aat Met Asn	gag to Glu Se: 430	gtg Val	att Ile	gtt Val	cga Arg 435	gct Ala	tcc Ser	gca Ala	ttg Leu	1348
att gcc Ile Ala 440	aga gca Arg Ala	atg aac Met Asn 445	gat at: Asp Ile	gtt Val	gga Gly	cat His 450	gaa Glu	gat Asp	gaa Glu	caa Gln	gaa Glu 455	1396
aga gga Arg Gly	cat gta His Val	gct tca Ala Ser 460	ctt att	gaa Glu	tgt Cys 465	tac Tyr	atg Met	aaa Lys	gat Asp	tat Tyr 470	g <b>ga</b> Gly	1444
gct tca Ala Ser	aag caa Lys Gln 475	gag act Glu Thr	tac att	aag Lys 480	ttc Phe	ctg Leu	aaa Lys	gag Glu	gtc Val 485	acc Thr	aat Asn	1492
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tta tat Leu Tyr 520	aag gag Lys Glu	aaa gat Lys Asp 525	aca tat Thr Ty	tca Ser	acc Thr	gcc Ala 530	aaa Lys	gga Gly	aaa Lys	ctt Leu	aaa Lys 535	1636
aac atg Asn Met	att aat Ile Asn	cca ata Pro Ile 540	cta att	gaa Glu	tct Ser 545	gtc Val	aaa Lys	ata Ile	taa *		·	1678
atataatg taaattga attgtttt tcatgttt ctttataa aataaaat	ag actca ta tttaa ta tctca tg tactg	acaaca to atttgc to atttcc ag gtaatc go	atgagttg aagccaaa gcttgtga tattgtat	gt taa ig tga ig tti it tgi	attoo aaata tggat tatto	tgg tac tat gtag	gate aaca tata tgtt	tttg acttg attgt gtc	gaa a gag t ta a	ataaa tgta attat	acaata attaaa ccatca	1738 1798 1858 1918 1978 2024

<211> 548

<212> PRT

<213> Lycopersicon esculentum

## <400> 52

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	485		90	4	195	
Ser Arg Pro	Thr Glu Val Pro 500	Met Phe V 505	al Leu Glu	Arg Val I	Leu Asn	
Leu Thr Arg 515	Val Ala Asp Thr	Leu Tyr L 520	ys Glu Lys	Asp Thr 'I 525	Tyr Ser	
Thr Ala Lys 530	Gly Lys Leu Lys 535	Asn Met I	le Asn Pro 540	Ile Leu I	lle Glu	
Ser Val Lys 545	Ile					
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<212>						
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		1	5	ile Asii I	ie Ala	
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10	Asn Ser Leu His 15	Asn Pne G	20	Pro Ser L	ys Ala 25	
tgg tct acc Trp Ser Thr	tct tgc act gca Ser Cys Thr Ala	ccc gca g Pro Ala A	ct cgc ctc	cgg gca t Arg Ala S	ct tcc Ser Ser	148
•	30		35	<b>.</b>	40	
tcc tta caa	caa gaa aaa cct Gln Glu Lys Pro	cac caa a	tc cga cgc	tct ggg g	gat tac	196
ser hed Gill	45	50 50	ie arg arg	ser Gly A	ap Tyr	
caa ccc tct	ctt tgg gat ttc	aat tac a	ta cag tct	ctc aac a	ict ccg	244
	Leu Trp Asp Phe				Chr Pro	
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Tyr Lys Glu	Gln Arg His Phe	Asn Arg G	ln Ala Glu	Leu Ile M	Met Gln	232
	80		85			
gtg agg atg Val Arg Met	ttg ctc aag gta Leu Leu Lys Val	aag atg g Lvs Met G	ag gca att : Slu Ala Ile :	caa cag t Gln Gln I	tg gag eu Glu	340
90	95	_, U	100	~~. 4	105	
	gac ttg caa tac					388
red ite Asp	Asp Leu Gln Tyr 110	-	eu Ser Tyr 15		In Asp .20	

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cac His	aat Asn	aat Asn 140	gat Asp	ttg Leu	tat Tyr	ttc Phe	aca Thr 145	gct Ala	ctt Leu	gga Gly	ttc Phe	aga Arg 150	atc Ile	ctc Leu	aga Arg		484
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gag Glu 170	aag Lys	tgc Cys	agt Ser	gat Asp	ttc Phe 175	aat Asn	gca Ala	aac Asn	ctt Leu	gct Ala 180	caa Gln	gat Asp	acg Thr	aag Lys	gga Gly 185		580
atg Met	tta Leu	caa Gln	ctt Leu	tat Tyr 190	gaa Glu	gca Ala	tct Ser	ttc Phe	ctt Leu 195	ttg Leu	aga Arg	gaa Glu	ggt Gly	gaa Glu 200	gat Asp		628
aca Thr	ttg Leu	gag Glu	cta Leu 205	gca Ala	aga Arg	cga Arg	ttt Phe	tcc Ser 210	acc Thr	aga Arg	tct Ser	cta Leu	cga Arg 215	gaa Glu	aaa Lys		676
Phe	Asp	Glu 220	Gly	Gly	Asp	gaa Glu	Ile 225	Asp	Glu	Asp	Leu	Ser 230	Ser	Trp	Ile	•	724
Arg	235	Ser	Leu	Asp	Leu	cct Pro 240	Leu	His	Trp	Arg	Val 245	Gln	Gly	Leu	Glu		772
250	Arg	Trp	Phe	Leu	Asp 255	gct Ala	Tyr	Ala	Arg	Arg 260	Pro	Asp	Met	Asn	Pro 265		820
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tat Tyr	caa Gln	gaa Glu	gaa Glu 285	ctg Leu	aaa Lys	gat Asp	atc Ile	tca Ser 290	agg Arg	tgg Trp	tgg Trp	aat Asn	agt Ser 295	tcg Ser	tgc Cys		916
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Lys 330	Met	Ala	Ala	Val	Ile 335	Ile	Thr	Phe	Ile	Thr 340	Ile	Ile	Asp	Asp	Val 345	
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gat Asp	att Ile 395	cta Leu	aaa Lys	gat Asp	caa Gln	cat His 400	ttc Phe	aac Asn	agċ Ser	atc Ile	cca Pro 405	tat Tyr	tta Leu	cag Gln	aga Arg	1252
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Gly	aca	tca	caa	cat	gag	ctg	gag Glu	aga Arq	gga Glv	gac Asp	gta Val	ccg Pro	aaa Lys	gca Ala	atc Ile	1540
490	1111	ser	GIII	птэ	495	204		J	•	500			•		505	
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cag Gln cac	tgc Cys gtg	tac	atg Met ttt	aat Asn 510	495 gac Asp	aca Thr	aat Asn gag	gct Ala gcg	tcg Ser 515	500 gag Glu aag	aga Arg gag	gag Glu atg	gcg Ala	gtg Val 520	505 gaa Glu gtc	1588 1636

gct aat ctt gca agg gcg gct cag ttt ata tat ctc gac ggg gat ggg Ala Asn Leu Ala Arg Ala Ala Gln Phe Ile Tyr Leu Asp Gly Asp Gly 555 560 565	1732
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ttc cag cct tat gtc tga ataaatcgaa aatccaacct actatgtatc Phe Gln Pro Tyr Val * 590	1828
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<211> 590

<212> PRT

<213> Salvia officinalis

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Leu His Trp Arg Val Gln Gly Leu Glu Ala Arg Trp Phe Leu Asp Ala
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                                     250
Tyr Ala Arg Arg Pro Asp Met Asn Pro Leu Ile Phe Lys Leu Ala Lys
                                265
Leu Asn Phe Asn Ile Val Gln Ala Thr Tyr Gln Glu Glu Leu Lys Asp
                            280
Ile Ser Arg Trp Trp Asn Ser Ser Cys Leu Ala Glu Lys Leu Pro Phe
                        295
                                             300
Val Arg Asp Arg Ile Val Glu Cys Phe Fhe Trp Ala Ile Ala Ala Phe
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                                        315
Glu Pro His Gln Tyr Ser Tyr Gln Arg Lys Met Ala Ala Val Ile Ile
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                                     330
Thr Phe Ile Thr Ile Ile Asp Asp Val Tyr Asp Val Tyr Gly Thr Ile
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Glu Glu Leu Glu Leu Leu Thr Asp Met Ile Arg Arg Trp Asp Asn Lys
                            360
Ser Ile Ser Gln Leu Pro Tyr Tyr Met Gln Val Cys Tyr Leu Ala Leu
                        375
                                             380
Tyr Asn Phe Val Ser Glu Arg Ala Tyr Asp Ile Leu Lys Asp Gln His
                    390
                                         395
Phe Asn Ser Ile Pro Tyr Leu Gln Arg Ser Trp Val Ser Leu Val Glu
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                                     410
Gly Tyr Leu Lys Glu Ala Tyr Trp Tyr Tyr Asn Gly Tyr Lys Pro Ser
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            420
                                                     430
Leu Glu Glu Tyr Leu Asn Asn Ala Lys Ile Ser Ile Ser Ala Pro Thr
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                                                 445
Ile Ile Ser Gln Leu Tyr Phe Thr Leu Ala Asn Ser Ile Asp Glu Thr
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Ala Ile Glu Ser Leu Tyr Gln Tyr His Asn Ile Leu Tyr Leu Ser Gly
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Thr Ile Leu Arg Leu Ala Asp Asp Leu Gly Thr Ser Gln His Glu Leu
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Glu Arg Gly Asp Val Pro Lys Ala Ile Gln Cys Tyr Met Asn Asp Thr
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                                 505
Asn Ala Ser Glu Arg Glu Ala Val Glu His Val Lys Phe Leu Ile Arg
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Glu Ala Trp Lys Glu Met Asn Thr Val Thr Thr Ala Ser Asp Cys Pro
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Phe Thr Asp Asp Leu Val Ala Ala Ala Ala Asn Leu Ala Arg Ala Ala
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                                         555
Gln Phe Ile Tyr Leu Asp Gly Asp Gly His Gly Val Gln His Ser Glu
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<sup>&</sup>lt;211> 2861 ~

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Abies grandis

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575

623

tgg ggt gaa gga ttc tac ttc ttg gca tat gac aga ata ctg gct aca

Trp Gly Glu Gly Phe Tyr Phe Leu Ala Tyr Asp Arg Ile Leu Ala Thr

ett gea tgt att att acc ett acc etc tgg egt act ggg gag aca caa

Leu Ala Cys Ile Ile Thr Leu Thr Leu Trp Arg Thr Gly Glu Thr Gln

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gta Val	cag Gln	aaa Lys 210	ggt Gly	att Ile	gaa Glu	ttc Phe	ttc Phe 215	agg Arg	aca Thr	caa Gln	gct Ala	gga Gly 220	aag Lys	atg Met	gaa Glu	671
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gca Ala 240	atg Met	cta Leu	aag Lys	gaa Glu	gct Ala 245	aaa Lys	atc Ile	tta Leu	ggc Gly	ttg Leu 250	gat Asp	ctg Leu	cct Pro	tac Tyr	gat Asp 255	767
ttg Leu	cca Pro	ttc Phe	ctg Leu	aaa Lys 260	caa Gln	atc Ile	atc Ile	gaa Glu	aag Lys 265	cgg Arg	gag Glu	gct Ala	aag Lys	ctt Leu 270	aaa Lys	815
agg Arg	att Ile	ccc Pro	act Thr 275	gat Asp	gtt Val	ctc Leu	tat Tyr	gcc Ala 280	ctt Leu	cca Pro	aca Thr	acg Thr	tta Leu 285	ttg Leu	tat Tyr	863
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ctt Leu	caa Gln 305	tcc Ser	aag Lys	gat Asp	gga Gly	tca Ser 310	ttt Phe	ctc Leu	agc Ser	tct Ser	ccg Pro 315	gca Ala	tct Ser	aca Thr	gcg Ala	959
gct Ala 320	gta Val	ttc Phe	atg Met	cgt Arg	aca Thr 325	ggg ggg	aac Asn	aaa Lys	aag Lys	tgc Cya 330	ttg Leu	gat Asp	ttc Phe	ttg Leu	aac Asn 335	1007
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gtt Val 400	cct Pro	gat Asp	att Ile	gat Asp	gat Asp 405	aca Thr	gcc Ala	atg Met	ggc Gly	ctt Leu 410	cga Arg	atc Ile	ttg Leu	aga Arg	tta Leu 415	1247

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aat Asn	gly aaa	gag Glu	ttc Phe 435	ttt Phe	Cys	ttc Phe	ttg Leu	ggt Gly 440	caa Gln	aca Thr	cag Gln	aga Arg	gga Gly 445	gtt Val	aca Thr		1343
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atg Met	cca Pro	agg Arg	ttg Leu 515	gag Glu	gct Ala	aga Arg	agc Ser	tat Tyr 520	att Ile	gaa Glu	aac Asn	tat Tyr	999 Gly 525	cca Pro	gat Asp	•	1583
gat Asp	gtg Val	tgg Trp 530	ctt Leu	gga Gly	aaa Lys	act Thr	gta Val 535	tat Tyr	atg Met	atg Met	cca Pro	tac Tyr 540	att Ile	tcg Ser	aat Asn		1631
gaa Glu	aag Lys 545	tat Tyr	tta Leu	gaa Glu	cta Leu	gcg Ala 550	aaa Lys	ctg Leu	gac Asp	ttc Phe	aat Asn 555	aag Lys	gtg Val	cag Gln	tc <b>t</b> Ser		1679
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ttc Phe	tca Ser	ccg Pro	gca Ala 595	tcc Ser	ttt Phe	atc Ile	ttt Phe	gag Glu 600	ccc Pro	gag Glu	ttt Phe	tct Ser	aag Lys 605	tgc Cys	aga Arg		1823
gag Glu	gtt Val	tat Tyr 610	aca Thr	aaa Lys	act Thr	tcc Ser	aat Asn 615	ttc Phe	act Thr	gtt Val	att Ile	tta Leu 620	gat Asp	gat Asp	ctt Leu		1871

			cat His														1919
			tgg Trp														1967
			ttt Phe														2015
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			gtc Val														2111
			tat Tyr													-	2159
			gca Ala													•	2207
ggg Gly	gag Glu	gtt Val	ctt Leu	aca Thr 740	gat Asp	gaa Glu	gta Val	ctc Leu	tcc Ser 745	aaa Lys	att Il <b>e</b>	gat Asp	cgc Arg	gaa Glu 750	tct Ser		2255
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			cag Gln														2351
caa Gln	tgt Cys 785	tat Tyr	atg Met	aag Lys	gac Asp	cat His 790	cct Pro	aaa Lys	atc Ile	tct Ser	gaa Glu 795	gaa Glu	gaa Glu	gct Ala	cta Leu		2399
			tat Tyr														2447
			aat Asn														2495
gaa	act	gca	aga	ata	atg	caa	ctc	ttt	tat	atg	caa	<b>9</b> 99	gat	ggt	ttg		2543

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Thr Leu Ser His Asp Met Glu Ile Lys Glu His Val Lys Asn Cys Leu
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                                                                      2639
Phe Gln Pro Val Ala
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Trp Gly Lys Gly Ser Asn Lys Ile Ile Ala Cys Val Gly Glu Gly Gly
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Glu Lys Arg Ile Glu Thr Leu Ile Ser Glu Ile Lys Asn Met Phe Arg
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                                                 125
Cys Met Gly Tyr Gly Glu Thr Asn Pro Ser Ala Tyr Asp Thr Ala Trp
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                                             140
Val Ala Arg Ile Pro Ala Val Asp Gly Ser Asp Asn Pro His Phe Pro
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                                         155
Glu Thr Val Glu Trp Ile Leu Gln Asn Gln Leu Lys Asp Gly Ser Trp
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                                     170
Gly Glu Gly Phe Tyr Phe Leu Ala Tyr Asp Arg Ile Leu Ala Thr Leu
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Ala Cys Ile Ile Thr Leu Thr Leu Trp Arg Thr Gly Glu Thr Gln Val
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                             200
Gln Lys Gly Ile Glu Phe Phe Arg Thr Gln Ala Gly Lys Met Glu Asp
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                         215
                                             220
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Ile Pro Thr Asp Val Leu Tyr Ala Leu Pro Thr Thr Leu Leu Tyr Ser
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                                                 285
Leu Glu Gly Leu Gln Glu Ile Val Asp Trp Gln Lys Ile Met Lys Leu
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                                             300
Gln Ser Lys Asp Gly Ser Phe Leu Ser Ser Pro Ala Ser Thr Ala Ala
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Val Phe Met Arg Thr Gly Asn Lys Lys Cys Leu Asp Phe Leu Asn Phe
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Val Leu Lys Lys Phe Gly Asn His Val Pro Cys His Tyr Pro Leu Asp
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                                 345
Leu Phe Glu Arg Leu Trp Ala Val Asp Thr Val Glu Arg Leu Gly Ile
                             360
Asp Arg His Phe Lys Glu Glu Ile Lys Glu Ala Leu Asp Tyr Val Tyr
                         375
Ser His Trp Asp Glu Arg Gly Ile Gly Trp Ala Arg Glu Asn Pro Val
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                                         395
Pro Asp Ile Asp Asp Thr Ala Met Gly Leu Arg Ile Leu Arg Leu His
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Gly Tyr Asn Val Ser Ser Asp Val Leu Lys Thr Phe Arg Asp Glu Asn
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                                         475
Leu Glu Asn Val Asp Ala Phe Asp Lys Trp Ala Phe Lys Lys Asn Ile
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Arg Gly Glu Val Glu Tyr Ala Leu Lys Tyr Pro Trp His Lys Ser Met
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Pro Arg Leu Glu Ala Arg Ser Tyr Ile Glu Asn Tyr Gly Pro Asp Asp
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                                                 525
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                                             540
Lys Tyr Leu Glu Leu Ala Lys Leu Asp Phe Asn Lys Val Gln Ser Ile
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Ser Pro Ala Ser Phe Ile Phe Glu Pro Glu Phe Ser Lys Cys Arg Glu
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Glu Val Leu Thr Asp Glu Val Leu Ser Lys Ile Asp Arg Glu Ser Arg
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Phe Leu Gln Leu Met Gly Leu Thr Gly Arg Leu Val Asn Asp Thr Lys
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Thr Tyr Gln Ala Glu Arg Gly Gln Gly Glu Val Ala Ser Ala Ile Gln
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Cys Tyr Met Lys Asp His Pro Lys Ile Ser Glu Glu Glu Ala Leu Gln
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His Val Tyr Ser Val Met Glu Asn Ala Leu Glu Glu Leu Asn Arg Glu
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Thr Ala Arg Ile Met Gln Leu Phe Tyr Met Gln Gly Asp Gly Leu Thr
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                                                                       111
              Met Ala Leu Leu Ser Ile Val Ser Leu Gln Val Pro Lys
tee tge ggg ctg aaa teg ttg ate agt tee age aat gtg cag aag get
                                                                       159
Ser Cys Gly Leu Lys Ser Leu Ile Ser Ser Ser Asn Val Gln Lys Ala
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ctc tgt atc tct aca gca gtc cca aca ctc aga atg cgt agg cga cag
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Leu Cys Ile Ser Thr Ala Val Pro Thr Leu Arg Met Arg Arg Gln
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Lys Ala Leu Val Ile Asn Met Lys Leu Thr Thr Val Ser His Arg Asp
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aac Asn	ctg Leu	tgg Trp 80	g <b>aa</b> Glu	gat Asp	gat Asp	ttc Phe	ata Ile 85	caa Gln	tca Ser	ttg Leu	tcc Ser	tca Ser 90	cct Pro	tat Tyr	Gly 999		351
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gat ( Asp																	735
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ccc Pro																	831
ttg Leu																	879

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			ctc Leu													1455
acc Thr	ttt Phe	atg Met	caa Gln 465	gaa Glu	gca Ala	aag Lys	tgg Trp	atc Ile 470	tcc Ser	agc Ser	agt Ser	tat Tyr	ctc Leu 475	cca Pro	acg Thr	1503

ttt Phe	gag Glu	gag Glu 480	tac Tyr	ttg Leu	aag Lys	aat Asn	gca Ala 485	aaa Lys	gtt Val	agt Ser	tct Ser	ggt Gly 490	tct Ser	cgc Arg	ata Ile		1551
gcc Ala	aca Thr 495	tta Leu	caa Gln	ccc Pro	att Ile	ctc Leu 500	act Thr	ttg Leu	gat Asp	gta Val	cca Pro 505	ctt Leu	cct Pro	gat Asp	tac Tyr		1599
ata Ile 510	ctg Leu	caa Gln	gaa Glu	att Ile	gat Asp 515	tat Tyr	cca Pro	tcc Ser	aga Arg	ttc Phe 520	aat Asn	gag Glu	tta Leu	gct Ala	tcg Ser 525		1647
tcc Ser	atc Ile	ctt Leu	cga Arg	cta Leu 530	cga Arg	ggt Gly	gac Asp	acg Thr	cgc Arg 535	Cya	tac Tyr	aag Lys	gcg Ala	gat Asp 540	agg Arg		1695
gcc Ala	cgt Arg	gga Gly	gaa Glu 545	gaa Glu	gct Ala	tca Ser	gct Ala	ata Ile 550	tcg Ser	tgt Cys	tat Tyr	atg Met	aaa Lys 555	gac Asp	cat His		1743
cct Pro	gga Gly	tca Ser 560	ata Ile	gag Glu	gaa Glu	gat Asp	gct Ala 565	ctc Leu	aat Asn	cat His	atc Ile	aac Asn 570	gcc Ala	atg Met	atc Ile	-	1791
agt Ser	gat Asp 575	gca Ala	atc Ile	aga Arg	gaa Glu	tta Leu 580	aat Asn	tgg Trp	gag Glu	ctt Leu	ctc Leu 585	aga Arg	ccg Pro	gat Asp	agc Ser		1839
aaa Lys 590	agt Ser	ccc Pro	atc Ile	tct Ser	tcc Ser 595	aag Lys	aaa Lys	cat His	gct Ala	ttt Phe 600	gac Asp	atc Ile	acc Thr	aga Arg	gct Ala 605		1887
ttc Phe	cat His	cat His	gtc Val	tac Tyr 610	aaa Lys	tat Tyr	cga Arg	gat Asp	ggt Gly 615	tac Tyr	act Thr	gtt Val	tcc Ser	aac Asn 620	aac Asn		1935
gaa Glu	aca Thr	aag Lys	aat Asn 625	ttg Leu	gtg Val	atg Met	aaa Lys	acc Thr 630	gtt Val	ctt Leu	gaa Glu	cct Pro	ctc Leu 635	gct Ala	ttg Leu		1983
taa *	aaa	catat	ag a	aatgo	catta	aa aa	atgto	gggaa	a gto	ctata	aatc	taga	actat	tc	,		2036

tctatctttc ataatgtaga tctggatgtg tattgaactc taaaaaaaaa aaa

<400> 58

Met Ala Leu Leu Ser Ile Val Ser Leu Gln Val Pro Lys Ser Cys Gly

<sup>&</sup>lt;210> 58 <211> 637

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Abies grandis

Leu Lys Ser Leu-Ile Ser Ser Ser Asn Val Gln Lys Ala Leu Cys Ile Ser Thr Ala Val Pro Thr Leu Arg Met Arg Arg Arg Gln Lys Ala Leu Val Ile Asn Met Lys Leu Thr Thr Val Ser His Arg Asp Asp Asn Gly Gly Gly Val Leu Gln Arg Arg Ile Ala Asp His His Pro Asn Leu Trp Glu Asp Asp Phe Ile Gln Ser Leu Ser Ser Pro Tyr Gly Gly Ser Ser Tyr Ser Glu Arg Ala Glu Thr Val Val Glu Glu Val Lys Glu Met Phe Asn Ser Ile Pro Asn Asn Arg Glu Leu Phe Gly Ser Gln Asn Asp Leu Leu Thr Arg Leu Trp Met Val Asp Ser Ile Glu Arg Leu Gly Ile Asp Arg His Phe Gln Asn Glu Ile Arg Val Ala Leu Asp Tyr Val Tyr Ser Tyr Trp Lys Glu Lys Glu Gly Ile Gly Cys Gly Arg Asp Ser Thr Phe Pro Asp Leu Asn Ser Thr Ala Leu Ala Leu Arg Thr Leu Arg Leu His Gly Tyr Asn Val Ser Ser Asp Val Leu Glu Tyr Phe Lys Asp Glu Lys Gly His Phe Ala Cys Pro Ala Ile Leu Thr Glu Gly Gln Ile Thr Arg Ser Val Leu Asn Leu Tyr Arg Ala Ser Leu Val Ala Phe Pro Gly Glu Lys Val Met Glu Glu Ala Glu Ile Phe Ser Ala Ser Tyr Leu Lys Lys Val Leu Gln Lys Ile Pro Val Ser Asn Leu Ser Gly Glu Ile Glu Tyr Val Leu Glu Tyr Gly Trp His Thr Asn Leu Pro Arg Leu Glu Ala Arg Asn Tyr Ile Glu Val Tyr Glu Gln Ser Gly Tyr Glu Ser Leu Asn Glu Met Pro Tyr Met Asn Met Lys Lys Leu Leu Gln Leu Ala Lys Leu Glu Phe Asn Ile Phe His Ser Leu Gln Leu Arg Glu Leu Gln Ser Ile Ser Arg Trp Trp Lys Glu Ser Gly Ser Ser Gln Leu Thr Phe Thr Arg His Arg His Val Glu Tyr Tyr Thr Met Ala Ser Cys Ile Ser Met Leu Pro Lys His Ser Ala Phe Arg Met Glu Phe Val Lys Val Cys His Leu Val Thr Val Leu Asp Asp Ile Tyr Asp Thr Phe Gly Thr Met Asn Glu Leu Gln Leu Phe Thr Asp Ala Ile Lys Arg Trp Asp Leu Ser Thr Thr Arg Trp Leu Pro Glu Tyr Met Lys Gly Val Tyr Met Asp Leu Tyr Gln Cys 

Ile Asn Glu Met Val Glu Glu Ala Glu Lys Thr Gln Gly Arg Asp Met Leu Asn Tyr Ile Gln Asn Ala Trp Glu Ala Leu Phe Asp Thr Phe Met Gln Glu Ala Lys Trp Ile Ser Ser Ser Tyr Leu Pro Thr Phe Glu Glu Tyr Leu Lys Asn Ala Lys Val Ser Ser Gly Ser Arg Ile Ala Thr Leu Gln Pro Ile Leu Thr Leu Asp Val Pro Leu Pro Asp Tyr Ile Leu Gln Glu Ile Asp Tyr Pro Ser Arg Phe Asn Glu Leu Ala Ser Ser Ile Leu Arg Leu Arg Gly Asp Thr Arg Cys Tyr Lys Ala Asp Arg Ala Arg Gly Glu Glu Ala Ser Ala Ile Ser Cys Tyr Met Lys Asp His Pro Gly Ser Ile Glu Glu Asp Ala Leu Asn His Ile Asn Ala Met Ile Ser Asp Ala Ile Arg Glu Leu Asn Trp Glu Leu Leu Arg Pro Asp Ser Lys Ser Pro Ile Ser Ser Lys Lys His Ala Phe Asp Ile Thr Arg Ala Phe His His Val Tyr Lys Tyr Arg Asp Gly Tyr Thr Val Ser Asn Asn Glu Thr Lys Asn Leu Val Met Lys Thr Val Leu Glu Pro Leu Ala Leu